

## DESCRIPTION

## PROBE SET AND METHOD FOR IDENTIFYING HLA ALLELE

## TECHNICAL FIELD

5           The present invention relates to a probe set and a method for identifying an allele of human HLA.

## BACKGROUND ART

          Human leukocyte antigen (HLA) is known to  
10   include multiple HLA types, such as HLA-A, HLA-B, HLA-C, HLA-DP, HLA-DQ, HLA-DR, and HLA-MICA. An HLA allele is designated with a four or more digit number by the WHO HLA Nomenclature Committee. The principle of the nomenclature is that the first two digits  
15   correspond to the serotypes; the third and fourth digits distinguish the alleles of different amino acid sequences (subtypes); and the fifth digit distinguishes the alleles of different base sequences but encoding the same amino acid sequence. Typing of  
20   these alleles has been conventionally conducted at the serological level. Although this serological method does not require special sample processing, and enables easy typing using antigen-antibody reaction, the serotypes are the roughest  
25   classification corresponding to the first two digits of the numbers according to the nomenclature described above.

Many of other commercially available kits of the type associated with genomic extraction do not have enough accuracy to identify each allele individually. It is the current state that such a  
5 kit distinguishes multiple alleles as a group. Moreover, even a kit based on the SBT (Sequencing Based Typing) method, which enables the most detailed polymorphic analysis, often fails to solve the problem of ambiguity by one analysis since most  
10 samples are heterozygotes requiring reexamination. Such problematic alleles are listed collectively in <http://www.ihwg.org/protocols/sbt/ambiguities2.pdf> by the International Histocompatibility Working Group (IHWG).

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#### DISCLOSURE OF INVENTION

On the other hand, with the development of advanced medical treatment in recent years, detailed HLA typing is required in organ transplantation, etc.  
20 In addition, associations of HLA with diabetes, cancer, and other multifactorial diseases have been suggested. With such a background, a test method is desired that can identify each allele individually. Upon such demands it is an object of the present  
25 invention to provide a probe set that is useful for identifying each allele of HLA individually, and a method for identification of an HLA allele by the use

thereof.

A probe set for identifying an allele of HLA according to the present invention is a probe set comprising multiple probes that can be used for  
5 identifying HLA allele contained in a specimen, characterized in that each of the multiple probes comprises a partial sequence containing a base represented by a capital letter in a sequence of each allele in an allele list in the description.

10 An embodiment of the present invention is a probe set comprising multiple probes that can be used for identification of an HLA-A allele in a specimen, characterized in that each of the multiple probes comprises a partial sequence containing a base  
15 represented by a capital letter in a sequence of each allele in an allele list in the description.

Another embodiment of the present invention is a method for identification of an HLA-A allele contained in a specimen using a probe set,  
20 characterized in that the probe set is the probe set described above.

Another embodiment of the present invention is a probe set comprising multiple probes that can be used for identification of an HLA-B allele in a  
25 specimen, characterized in that each of the multiple probes comprises a partial sequence containing a base represented by a capital letter in a sequence of each

allele in an allele list in the description.

Another embodiment of the present invention is a method for identification of an HLA-B allele contained in a specimen using a probe set,

5 characterized in that the probe set is the probe set described above.

Another embodiment of the present invention is a probe set comprising multiple probes that can be used for identification of an HLA-C allele in a  
10 specimen, characterized in that each of the multiple probes comprises a partial sequence containing a base represented by a capital letter in a sequence of each allele in an allele list in the description.

Another embodiment of the present invention is  
15 a method for identification of an HLA-C allele contained in a specimen using a probe set, characterized in that the probe set is the probe set described above.

Another embodiment of the present invention is  
20 a probe set comprising multiple probes that can be used for identification of an HLA-DP allele in a specimen, characterized in that each of the multiple probes comprises a partial sequence including a base represented by a capital letter in the sequence of  
25 each allele in the allele list in the description.

Another embodiment of the present invention is a method for identification of an HLA-DP allele



contained in a specimen using a probe set,  
characterized in that the probe set is the probe set  
described above. Another embodiment according to the  
present invention is a probe set for identification  
5 of an HLA-DQ allele that is a probe set comprising  
multiple probes that can be used for identification  
of an HLA-DQ allele in a specimen, characterized in  
that each of the multiple probes comprises a partial  
sequence containing a base represented by a capital  
10 letter in a sequence of each allele in an allele list  
in the description.

Another embodiment of the present invention is  
a probe set comprising multiple probes that can be  
used for identification of an HLA-DQ allele in a  
15 specimen, characterized in that each of the multiple  
probes comprises a partial sequence containing a base  
represented by a capital letter in a sequence of each  
allele in an allele list in the description.

Another embodiment of the present invention is  
20 a method for identification of an HLA-DQ allele  
contained in a specimen using a probe set,  
characterized in that the probe set is the probe set  
described above.

Another embodiment of the present invention is  
25 a probe set comprising multiple probes that can be  
used for identification of an HLA-DR allele in a  
specimen, characterized in that each of the multiple

probes comprises a partial sequence containing a base represented by a capital letter in a sequence of each allele in an allele list in the description.

Another embodiment of the present invention is  
5 a method for identification of an HLA-DR allele contained in a specimen using a probe set, characterized in that the probe set is the probe set described above.

Another embodiment of the present invention is  
10 a probe set comprising multiple probes that can be used for identification of an HLA-MICA allele in a specimen, characterized in that each of the multiple probes comprises a partial sequence containing a base represented by a capital letter in a sequence of each  
15 allele in an allele list in the description.

Another embodiment of the present invention is a method for identification of an HLA-MICA allele contained in a specimen using a probe set, characterized in that the probe set is the probe set  
20 described above.

The probe set according to the present invention, and identification of an allele of each HLA type by the use thereof can contribute to diathesis diagnoses and tailor-made medicines, which  
25 are required in organ transplantation, cancer, diabetes, and other multifactorial diseases.

Other features and advantages of the present

invention will be apparent from the following description.

#### BEST MODE FOR CARRYING OUT THE INVENTION

5 Preferred embodiments of the present invention will now be described in detail. Each probe that constitutes the probe set of the present invention has a partial sequence including a base represented by a capital letter in each allele sequence in the  
10 allele lists described later. Preferably, segments consisting of 10 to 30 bases including a base represented by a capital letter are selected from each allele sequence, and the probe set is composed of probes having the obtained partial base sequences  
15 respectively. As specific examples, the following compositions can be employed:

1) A probe set for HLA-A allele identification consisting of respective probes listed in one of the probe list A1 shown in Tables 1-1 to 1-7 and the  
20 probe list A2 shown in Tables 2-1 to 2-6 shown later;

2) A probe set for HLA-B allele identification consisting of probes listed in one of the probe list B1 shown in Tables 5-1 to 5-9 and the probe list B2 shown in Tables 6-1 to 6-8 shown later;

25 3) A probe set for HLA-C allele identification consisting of probes listed in one of the probe list C1 shown in Tables 9 and the probe list C2 shown in

Table 10 shown later;

4) A probe set for HLA-DP allele identification consisting of probes listed in one of the probe lists DP1-DP4 shown in Tables 13-1 to 16-5 respectively as  
5 shown later;

5) A probe set for HLA-DQ allele identification consisting of probes listed in one of the probe lists DQ1 to DQ 4 shown in Tables 17A, 17B-1, 17B-2, 18A, 18B-1 and 18B-2 respectively as shown later;

10 6) A probe set for HLA-DR allele identification consisting of probes listed in one of the probe lists DR1 and DR2 shown in Tables 21-1 to 21-8 and Tables 22-1 to 22-7 respectively, as shown later; and

7) A probe set for HLA-MICA allele  
15 identification consisting of probes listed in one of the probe lists MICA1 and MICA2 shown in Tables 25-1, 25-2 and Tables 26-1 to 26-2 respectively, as shown later.

For example, the No.0 probe in the probe list  
20 A1 has a 16-base sequence of "gccccgcttcacgcC", which is a segment containing the first capital lettered base C in A\*010101, and the No.0 probe in the probe list 2 has an 18-base sequence of "cttcacgcCgtgggcta", which is a segment also  
25 containing the first capital lettered base C in the same allele.

In the allele list, each allele is assigned

with a unique number such as "A\*xxxx" in accordance with "allele nomenclature" by Japanese Society for Histocompatibility and Immunogenetics, HLA Standardization Committee.

5           To identify an allele using a probe set according to the present invention, two methods are possible: one is detection by hybridization; and the other is direct detection by PCR without hybridization. In either method, each probe is an  
10 oligonucleotide of preferably more than 10 and less than 30 nucleotides in length and designed to include the base represented by a capital letter, i.e., a base specific for the allele to be identified.

          Moreover, the probe arrays provided in the  
15 present invention present groups of varied bases for identification of each allele individually by positions chosen for the probes. As a method for detection of such a varied base, the detection method by hybridization, and the method of direct detection  
20 by PCR without hybridization can also be preferably used. Also in these cases, the probes are designed as oligonucleotides of preferably more than 10 and less than 30 nucleotides in length each containing a base represented by a capital letter.

25           When a variation is detected by hybridization, probes are preferably designed to have a variant base represented by a capital letter near the center of

the probes, which makes  $T_m$  difference between full-matched and mismatched pairs larger, enabling easier separation of them by adjusting the reaction temperature of hybridization.

5           On the other hand, when the variation is directly detected by PCR, the variant base is rather placed near the 3' end so that enzymatic recognition and elongation of annealed double strands will not occur. Also, some variation methods are possible,  
10 such as a method placing a variant base at the second from the 3' end an artificial variant base at the third from the 3' end as with Allele Specific Primer (Toyobo Co., Ltd.); a method circularizing probes by ligation with a mismatch placed near the 3' end  
15 (Amersham Biosciences Co., Ltd.); TaqMan-MGB (ABI Co.); and 3'-end mismatch using LNA (Proligo Japan Co., Ltd.).

For example, a segment including the fourth capital letter of A\*2302 is "ggagcagTggagagC", and  
20 the corresponding segment of A\*2303 of the same serotype is "ggagcagtTgagagc", differing at the ninth base. By using a probe with a sequence of one of these segments, one can be distinguished from the other by mismatching.

[EXAMPLES]

The present invention will be described further by way of examples in the following. The SEQ ID NOS of the sequences in Examples are in the following

5 relation:

Examples 1 and 2: Sequence Listing A

Examples 3 and 4: Sequence Listing B

Examples 5 and 6: Sequence Listing C

Examples 7 and 8: Sequence Listing DP

10 Examples 9 and 10: Sequence Listing DQ

Examples 11 and 12: Sequence Listing DR

Examples 13 and 14: Sequence Listing MICA.

(Example 1)

15 Probes for identification of HLA-A allele

Extraction of DNA from 1 ml of human blood was performed using GFX Genomic Blood DNA Purification Kit from Amersham Biosciences. The protocol is as follows:

20 Blood 1 ml →

Add RBC Lysis Solution [hemolysate] →

Mix gently at room temperature for 5 minutes →

Centrifuge at 12,000-16,000 × g for 20 seconds →

Discard the supernatant leaving 20-50 µl →

25 Resuspend the precipitation →

Add Extraction Solution and vortex vigorously →

Stand at room temperature for 5 minutes [extraction

of DNA] →

Set a GFX Column in a Collection Tube →

Heat the elution buffer to 70°C →

Add the sample →

- 5 Centrifuge at 5,000 × g for 1 minute (binding of DNA)

→

Add Extraction Solution (washing) →

Centrifuge at 5,000 × g for 1 minute →

Add Washing Solution (washing) →

- 10 Centrifuge at 12,000 × g for 3 minutes →

Set a GFX Column in a centrifugal tube →

Eluate with pure water →

Stand at room temperature for 1 minute →

Centrifuge at 5,000-8,000 × g for 1 minute →

- 15 Concentrate to 230 µl ... .. solution (1).

Next, quantitative PCR was carried out using  
QuantiTect SYBR Green PCR Kit from QIAGEN and  
GeneAmp5700 from ABI. The reaction composition and  
20 the protocol are shown below.

- 1) Reaction composition/well (96 well microplate)

QuantiTect SYBR Green 2 × premix: 10 µl

Solution (1): 1 µl

Solution of one of the probes in the probe list

- 25 A1(10 pmol/µl): 1 µl

Mixed primers (10 pmol/µl)\*: 3 µl

Ultra pure water: 5 µl



(Total: 20  $\mu$ l)

\*consisting of 1  $\mu$ l each of the solutions respectively containing probes of the following sequences at 10 pmol/ $\mu$ l:

5 CCCATCTCAGGGTGAGGGGCT (SEQ ID NO: 632)

GCGCTGCAGCGTCTCCTTCC (SEQ ID NO: 633)

GCCCAGGTCTGGGTCAGGGCCAG (SEQ ID NO: 634)

2) PCR program

94°C: 180 sec followed by 30 cycles of [94°C: 10  
10 sec  $\rightarrow$  66°C: 10 sec  $\rightarrow$  72°C: 20 sec.].

Referring to Amp Plot and Dissociation curves on a display of 5700 software, and to the allele-probe correspondence list A1 (Tables 3-1 to 3-9), it was identified as A\*2402101.

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(Example 2)

Extraction of DNA from 1 ml of human blood was performed in the same manner as in Example 1. PCR of human HLA-A was then performed using ABI 9700 PCR  
20 Instrument and Ex Taq from Takara Bio Inc. The reaction composition and the protocol are as follows:

1) Reaction composition/tube

Ex Taq 2  $\times$  premix: 20  $\mu$ l

Solution (1): 3  $\mu$ l

25 Cy-3 dUTP (1 mM): 2  $\mu$ l

Mix primer (10 pmol/ $\mu$ l)\*: 3  $\mu$ l

Ultra pure water: 12  $\mu$ l

(Total: 40  $\mu$ l)

\*consisting 1  $\mu$ l each of the solutions respectively containing probes of the following sequences at 10 pmol/ $\mu$ l:

- 5           ATGGCTCCCCGAACCCTC (SEQ ID NO: 635)  
            ATGGCGCCCCGAACCCTC (SEQ ID NO: 636)  
            CATCTCAGGGTGAGGGGCT (SEQ ID NO: 637)

2) PCR program

94°C: 180 sec followed by 30 cycles of [94°C: 10  
10 sec → 66°C: 10 sec → 72°C: 20 sec]

After the completion of the reaction, unreacted dNTPs, etc., were removed using a purification column (QIAGEN QIAquick PCR Purification Kit) to obtain a sample.

- 15           At the same time, a DNA microarray was prepared to identify the allele in the specimen described above. The method for the preparation was in accordance with examples in Japanese Patent Application Laid-Open No. H11-187900. SH group was  
20 used as the functional group for immobilization. A glass substrate was treated by a silane-coupling agent to bind the SH group of the probes via a divalent reagent EMCS (N-(6-maleimidocaproyloxy)succinimide). Each probe in the  
25 probe list A2 was used for each dot.

The DNA microarray was blocked in advance with PBS supplemented with 1 wt% of BSA (bovine serum

albumin) for two hours. The sample was adjusted to have a salt concentration equal to that of the PBS, and to contain 0.1 wt% of SDS (sodium dodecyl sulfate) and 25% of formamide.

5           Then, hybridization was performed using the above sample (PCR product) and the prepared DNA microarray. 50  $\mu$ l of the sample was reacted with the blocked DNA microarray at 60 °C for 2 hours. Unreacted substances were washed off by washing three  
10 times with 2  $\times$  SSC solution (NaCl 300 mM, Sodium Citrate (trisodium citrate dihydrate,  $C_6H_5Na_3 \cdot 2H_2O$ ) 30 mM, pH 7.0), followed by washing twice with 0.1  $\times$  SSC solution. The DNA microarray was air-dried and the fluorometry measurement was conducted using  
15 GenePix4000B made by Axon. Referring to the allele-probe list A2 (Tables 4-1 to 4-9), the sample was identified as A\*2402101.

A\*010101 :

atggcgcgcatggcgccccgaacccctcctctgctactctcgggggccctggccctgaccagacctgggcgggct  
20 cccactccatgagglatttcttcacatccgigtgccggcccgccgaggagccccgcttcatcgCgtgggcta  
cgtggacgacacgcagttcgtgcggttcgacagcgacgccgagccagaagatggagccgcccggcgccgtggata  
gagcaggagggggccggagtattgggaccaggagacacggaatAtgaaggccctcacagactgaccgagcgaacc  
tggggacccctgcgcggctactacaaccagagcgaggacggttctcacacatccagataatgtatggctgcgacgt  
ggggccggacggcgcttccctccgcggtaccggcaggacgcctacgacggcaaggattacatcgccctgaacgag  
25 gacctgcgctcttggaccgcgccggacatggcagctcagatcaccaagcgcaagtgggaggcggtccatcgccggg  
agcagcggagagTctacctggagggcCGgtgcgtggacgggctccgcagataacctggagaacgggaaggagacgct  
gcagcgacggaccccccaagacacataatgaccaccaccccatctctgacatgaggccacctgagggtgcctgg

gcccTgGcttTtaccctgcggagatcacactgacctggcagcgggTtggggaggaccagaccaggacacggagc  
tcgtggagaccaggccTgcaggggatggaacctTccagaagtgggcggctgtggTggtGcctTtTggaGaggagca  
gagatacacctgccatgtgcagcatgagggTctgcccagccccTcacctgagatgggag (SEQ ID  
NO:1) ;

5 A\*010102 :

gctccactccatgaggtattTctTcacatccgtgtcccggccccggccgaggagccccgtTcatcgccgtggg  
ctactgtggacgacacgcagTtcgtgcggTtcgacagcgacgccgcgagccagaagtggagccgaggcgccgtgg  
atagagcaggagggggccggagtattgggaccaggagacacggaatatgaaggccactcacagactgaccgagcga  
acctggggacccTgcgcggctactacaaccagagcgaggacggTtTcacaccatccagataatgtatggctgcga  
10 cgtggggccggacggcgctTctccgcgggtaccggcaggacgcctacgacggcaaggattacatgccccTgaac  
gaggacctgcgtctTtgaccgcggcgacatggcagctcagatTaccaagcgcaagtgggaggcggtccatgcgg  
cggagcagcggagagTtacctggagggccggTgcgtggacggctccgcagatactggagaacgggaaggagac  
gctgcagcgacgg (SEQ ID NO: 2) ;

A\*0102 :

15 atggccgtcatggcggcccgaaacctTctTctgtactTctggggggccctggccctgaccagacctgggcgggct  
cccactccatgaggtattTctTcacatccgtgtcccggccccggcagTggAgagccccgtTcatcgcagTgggcta  
cgtggacgacacgcagTtcgtgcggTtcgacagcgacgccgcgagccagaagtggagccgaggcgccgtggata  
gagcaggagggggccggagtattgggaccaggagacacggaatatgaaggccactcacagactgaccgagcgaacc  
tggggacccTgcgcggctactacaaccagagcgaggacggTtTcacaccatccagataatgtatggctgcgacgt  
20 gggggccggacggcgctTctccgcgggtaccggcaggacgcctacgacggcaaggattacatgccccTgaacgag  
gacctgcgtctTtgaccgcggcgacatggcagctcagatcaccaagcgcaagtgggaggcggtccatgcggcgg  
agcagcggagagTtacctggagggccggTgcgtggacggctccgcagatactggagaacgggaaggagacgt  
gcagcgacggaccccccaagacacataTgaccaccaccccaTctTgacctgaggccacctgaggtgctgg  
gccctgggctTtTaccctgcggagatcacactgacctggcagcgggatggggaggaccagaccaggacacggagc  
25 tcgtggagaccaggccTgcaggggatggaacctTccagaagtgggcggctgtggTggtgcctTtTggagaggagca  
gagatacacctgccatgtgcagcatgagggTctgcccagccccTcacctgagatgggag (SEQ ID NO: 3)

A\*0103 :

gctcccactccatgaggtatttcttcacatccgtgtcccggcccgccgaggagccccgcttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagccagaagaaggagccgaggcgccgtgg  
atagagcaggaggggcccggagtattgggaccaggagacacggaatatgaagggccactcacagactgaccgagcga  
acctggggacccctgcgcggctactacaaccagagcgaggacggttctcacaccatccagatGatgtatggctgcga  
5 cgtggggccggacgggcgcttctccgcggtaccggcaggacgcctacgacggcaaggattacatcgccctgaac  
gaggacctgcgtcttggaccgcgccggacatggcagctcagatcaccaagcgcaagtgggaggcggtccatgcgg  
cggagcagcgagagcttacctggagggccggctgcgtggacgggctccgcagatacctggagaacgggaaggagac  
gctgcagcgacgg (SEQ ID NO: 4) ;

A\*0106 :

10 gctcccactccatgaggtatttcttcacatccgtgtcccggcccgccgaggagccccgcttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagccagaagaaggagccgaggcgccgtgg  
atagagcaggaggggcccggagtattgggaccaggagacacggaatatgaagggccactcacagactgaccgagcga  
acctggggacccctgcgcggctactacaaccagagcgaggacggttctcacaccatccagataatgtatggctgcga  
cgtggggccggacgggcgcttctccgcggtaccggcaggacgcctacgacggcaaggattacatcgccctgaac  
15 gaggacctgcgtcttggaccgcgccggacatggcagctcagatcaccaagcgcaagtgggaggcggtccatgcgg  
cggagcagTTgagagcctacctggagggccggctgcgtggacgggctccgcagatacctggagaacgggaaggagac  
gctgcagcgacgg (SEQ ID NO:5) ;

A\*0107 :

gctcccactccatgaggtatttcttcacatccgtgtcccggcccgccgaggagccccgcttcatcgccgtggg  
20 ctacgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagccagaagaaggagccgaggcgccgtgg  
atagagcaggagaggccTgagtattgggaccaggagacacggaatgtgaagggccactcacagactgaccgagAga  
acctggggacccctgcgcggctactacaaccagagcgaggCgggttctcacaccatccagataatgtatggctgcga  
cgtggggccggacgggcgcttctccgcggtaccggcaggacgcctacgacggcaaggattacatcgccctgaac  
gaggacctgcgtcttggaccgcgccggacatggcagctcagatcaccaagcgcaagtgggaggcggtccatgcgg  
25 cggagcagcgagagcttacctggagggccggctgcgtggacgggctccgcagatacctggagaacgggaaggagac  
gctgcagcgacgg (SEQ ID NO: 6) ;

A\*0108 :

gctcccactccatgaggtatttcttcacatccgtgtcccggcccgccgaggagccccgtttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagccagaagatggagccgaggcgccgtgg  
atagagcaggaggggcccggagtattgggaccaggagacacggaatatgaaggccactcacagactgaccgagcga  
acctggggacccctgcgcggctactacaaccagagcgaggacgggttcacaccatccagataatgtatggctgcga  
5 cgiggggccggacgggcgcttctccgcgggtaccggcaggacgcctacgacggcaaggattacatcgccctgaac  
gaggacctgcgctcttggaccgcggcgacatggcagctcagatcaccaagcgcaagtgggaggcggtccatgcgg  
cggagcagcggagagttacctggagggcTgggtgcgtggacgggtccgcagataacctggagaacgggaaggagac  
gctgcagcgacgg (SEQ ID NO: 7) ;

A\*0109 :

10 atggccgtcatggcggcccggaaccttcttctgtactctcggggccciggccctgacctgagacctgggcgggct  
cccactccatgaggtatttcttcacatccgtgtcccggcccgccgaggagccccgtttcatcgccgtgggcta  
cgtggacgacacgcagttAggtgcggttcgacagcgacgccgcgagccagaagatggagccgaggcgccgtggata  
gagcaggaggggcccggagtattgggaccaggagacacggaatatgaaggccactcacagactgaccgagcgaacc  
tggggacccctgcgcggctactacaaccagagcgaggacgggttcacaccatccagataatgtatggctgcgacgt  
15 ggggcccggacgggcgcttctccgcgggtaccggcaggacgcctacgacggcaaggattacatcgccctgaacgag  
gacctgcgctcttggaccgcggcgacatggcagctcagatcaccaagcgcaagtgggaggcggtccatgcggcgg  
agcagcggagagttacctggagggccgggtgcgtggacgggtccgcagataacctggagaacgggaaggagacgct  
gcagcgacggaccccccaagacacatatgaccaccacccatctctgacctgaggccacctgagggtcgtgg  
gcccgtgggtttctacctgcggagatcacactgacctggcagcgggatggggaggaccagaccaggacacggagc  
20 tctgtggagaccaggccctgcaggggatggaaccttccagaagtgggcccgtgtgtgtgtgccttctggagaggagca  
gagatacacctgccatgtgcagcatgagggtctgcccagccccctacccctgagatgggag (SEQ ID NO:  
8) ;

A\*020101 :

atggccgtcatggcggcccggaaccttcttctgtactctcggggcTciggccctgacctgagacctgggcgggct  
25 ctactccatgaggtatttcttcacatccgtgtcccggcccgccgaggagccccgtttcatcgcagttgggcta  
cgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagccagaggatggagccgaggcgccgtggata  
gagcaggaggggtccggagtattgggacggggagacacggaaAgitgaaggccactcacagactCaccgagttggacc

- tggggacctgctgggtactacaaccagagcgaggccggttctcacaccGtccagaGgatgtatggctgctgacgt  
ggggctggacTggcgcttctccgcggtaccaccagtagcctacgacggcaaggattacatcgccctgaaagag  
gacctgctcttggaccgcgcgacatggcagctcagaccaccaagcacaagtgggaggcgcccatgTggcgg  
agcagtTgagagcctacctggagggaCgtgctggagtggctccgcagataacctggagaacgggaaggagacgt  
5 gcagcgacggacgccccaaaacgcatatgacacaccagctgtctctgacctgaagccacctgaggtgctgg  
gacctgagcttctacctgcgagatcacactgacctggcagcggaaggaggaccagaccaggacacggagc  
tctggagaccaggctgcagggatggaaccttccagaagtggcggtgtgtgtggcttctggacaggagca  
gagataacctgccatgtgcagcatgagggtTgcccagccccctacctgagatgggag (SEQ ID NO:  
9) ;
- 10 A\*020102 :  
gctccacctccatgaggtatttcttcacatccgtgtccggcccgccgaggagccccgttcaicgcAgtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgagccagaggatggagccgcgggcgccgtgg  
atagagcaggagggtccgagtatgggacgGggagacacggaaAgigaaggccacctcacagactCaccgagtgg  
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5 A\*0213 :

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5 A\*0216 :

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5 A\*0221 :

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A\*0222 :

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15 A\*0225 :  
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10 A\*0230 :

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A\*0233 :

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NO:48) ;

20 A\*0234 :

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A\*0235 :

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15 A\*0236 :

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A\*0237 :

25 gctctcactccatgagglattttcttcacatccgtgtcccgccccggcgcggggagccccgcttcatcgcatggg  
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A\*0238 :  
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15 A\*0239 :  
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A\*0240 :  
25 gctctcactccatgaggatattcttcacatccgtgtcccgccccggccgcggggagccccgcttcatcgcagtggg  
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- acctggggaccctgcgcggctactacaaccagagcgaggccggttctcacaccgtccagaggatgtatggctgcga  
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5 gctgcagcgcacgg (SEQ ID NO: 55) ;  
A\*0241 :  
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10 acctggggaccctgcgcggctactacaaccagagcgaggccggttctcacaccGtccagaGgatgtatggctgcga  
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15 A\*0242 :  
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20 cgtgggggtcggactggcgcttccctcgcgggtaccaccagtacgcctacgacggcaaggattacatcgccctgaaa  
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A\*0244 :  
25 gctctcactccatgaggtaattctacacCtccgtgtcccggcccgccgaggagccccgcttcatcgcagtggg  
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A\*0245 :

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10 accitggggacccitgcgcggctactacaaccagagcgaggccgggttctcacaccGtccagaGgatgtaiggctgcga  
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15 A\*0246 :

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25 A\*0247 :

gctctcactccatgaggtatttcttcacatccgtgtcccgccccggccgcggggagccccgcttcatcgagltggg  
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A\*0248 :

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A\*0249 :

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25 A\*0250 :

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A\*0251 :

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A\*0252 :

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25 gctgcagcgcacgg (SEQ ID NO:66) ;

25 A\*0254 :

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5 cggagcagcAgagagcctacctggagggcaCgtgcgtggacgggctccgcagataacctggagaacgggaaggagac  
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A\*0255 :

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15 gctgcagcgcacgg (SEQ ID NO:68) ;

A\*0256 :

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A\*0257 :

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5 cgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagccagaggatggagccgcggggcgccgtggata  
gagcaggaggggtccggaglatgggacggggagacacggaaagtgaaggccactcacagactcaccgagtgacc  
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tctggagaccaggcctgcagggatggaaccttcagaagtggcggtgttggtggcttctggacaggagca  
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15 NO:70) ;

A\*0258 :

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25 A\*0259 :

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A\*0304 :

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25 NO:78) ;

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A\*0306 :

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A\*0308 :

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A\*0309 :

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A\*0310 :

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A\*110102 :

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A\*1103

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25 A\*1104 :

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NO:89) ;  
A\*1105 :  
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25 NO:90) ;  
A\*1106 :  
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A\*1107 :

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A\*1108 :

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A\*1109 :

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A\*1110 :

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A\*1111 :

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A\*1112 :

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A\*1113 :

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A\*1114 :

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A\*2302 :

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A\*2303 :

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A\*2305 :

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A\*2306 :

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A\*240203 :

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NO:113) ;

A\*2405 :

gctcccatccatgaggtatttctccacatccgtgtcccgcccgccgaggagccccgcttcatcgccgtggg  
5 ctacgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagccagaggatggagccgaggcgccgtgg  
atagagcaggagggcgaggatattgggacgaggagacaGggaaagtgaaggccactcacagactgaccgagaga  
acctgcggatcgcgctccgctactacaaccagagcgaggccggttctcacacctccagatgatgttggctgcga  
cgtggggtcggacggcgcttccctccgcggtaccaccagttacgctacgacggcaaggattacatcgccctgaaa  
gaggacctgcgctcttggaccgcgccgagacatggcggtcagatcaccCagcgcaagtgggaggcgcccatgtgg  
10 cggagcagcAgagagCctacctggagggaCgtgcgtggacgggtccgcagatacctggagaacgggaaggagac  
gtgcagcgacgg (SEQ ID NO:114) ;

A\*2406 :

gctcccatccatgaggtatttctccacatccgtgtcccgcccgccgaggagccccgcttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagccagaggatggagccgaggcgccgtgg  
15 atagagcaggagggcgaggatattgggacgaggagacaGggaaagtgaaggccactcacagactgaccgagaga  
acctgcggatcgcgctccgctactacaaccagagcgaggccggttctcacacctccagatgatgttggctgcga  
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gaggacctgcgctcttggaccgcgccgagacatggcggtcagatcaccaagcgcaagtgggaggcgcccatgtgg  
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20 gctgcagcgacgg (SEQ ID NO:115) ;

A\*2407 :

atggccgtcatggcgccccgaacctcgctctgtactctcgggggccctggccctgaccagacctgggcaggct  
cccactccatgaggtatttctccacatccgtgtcccgcccgccgaggagccccgcttcatcgccgtgggcta  
cgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagccagaggatggagccgaggcgccgtggata  
25 gacgaggagggcgaggatattgggacgaggagacagggaaagtgaaggccaGtcacagactgaccgagagaacc  
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5 ttgtggagaccaggcctgcaggggatggaaccttccagaagtgggcAgctgtgggtgtaccttctggagaggagca  
gagatacacctgccatgtgcagcatgagggctgcccgaagccctcacctgagatgggag (SEQ ID  
NO:116) ;  
A\*2408 :  
atggccgctatggcggccgaacctcgtccgtctactctcggggggccctggccctgaccagacctgggcaggct  
10 cccaAtccatgaggtatcttccacatccgtgtcccgcccgccggcgaggagccccgcttcatcgccgtgggcta  
cgtggacgacacgcagttcgtgcggttcgacagcgacccgcgagccagaggatggagccgcccggcgccgtggata  
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15 gacctgcgctcttggaccgcggcggacatggcggctcagatcaccaagcgcaagtgaggaggcgcccatgtggcgg  
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gcagcgcacggaccccccaagacacataatgaccaccaccccatctctgacctgaggccactctgagatgctgg  
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ttgtggagaccaggcctgcaggggatggaaccttccagaagtgggcagctgtgggtgtAccttctggagaggagca  
20 gagatacacctgccatgtgcagcatgagggctgcccgaagccctcacctgagatgggag (SEQ ID  
NO:117) ;  
A\*2410 :  
gtccccactccatgaggtatcttccacatccgtgtcccgcccgccggcgaggagccccgcttcatcgccgtggg  
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25 atagagcaggagggggccggagtattgggacgaggagacaGggaaagtgaaggccctcacagactgaccgagaga  
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cgtggggtcgacgggcgttccctccggggtaccaccagtagcctacgacggcaaggattacatgccccgaaa

gaggacctgcgctcttggaccgcgccggacatggcggctcagatcaccaagcgcaagtgggaggcgcccatgtgg  
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gctgcagcgcacgg (SEQ ID NO:118) ;

A\*2413 :

5 gctcccactccatgaggtatttctccacatccgtgtcccggcccgccgaggagccccgcttcatcgccgtggg  
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acctgcggatcgcgctccgctactacaaccagagcgaggccggttctcacacctccagatgaigtitggctgcga  
cgtggggctggacggcgcttccctccgcggtaccaccagtacgcttacgacggcaaggattacatcgccctgaaA  
10 gaggacctgcgctcttggaccgcgccggacatggcGgctcagatcaccaagcgcaagtgggaggcgcccatgTgg  
cggagcagcTgagagCctacctggaggccggtgcgtggacgggctccgcagataacctggagaacgggaaggagac  
gctgcagcgcacgg (SEQ ID NO:119) ;

A\*2414 :

gctcccactccatgaggtatttctccacatccgtgtcccggcccgccgaggagccccgcttcatcgccgtggg  
15 ctacgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagccagaggatggagccgcccgcgcgtgg  
atagagcaggagggccggagiatitgggacgaggagacaGggaaagigaagcccactcacagactgaccgagaga  
acctgcggatcgcgctccgctactacaaccagagcgaggccggttctcacaccgtccagaggatgtaigtgctgcga  
cgtggggctggacTggcgcttccctccgcggtaccaccagTaccgcttacgacggcaaggattacatcgccctgaaA  
gaggacctgcgctcttggaccgcgccggacatggcGgctcagatcaccaagcgcaagtgggaggcgcccatgtgg  
20 cggagcagcAgagagCctacctggaggccggtgcgtggacgggctccgcagataacctggagaacgggaaggagac  
gctgcagcgcacgg (SEQ ID NO:120) ;

A\*2415 :

gctcccactccatgaggtatttctccacatccgtgtcccggcccgccgaggagccccgcttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagccagaggatggagccgcccgcgcgtgg  
25 atagagcaggagggccggagiatitgggacgaggagacaGggaaagigaagcccactcacagactgaccgagaga  
acctgcggatcgcgctcGctactacaaccagagcgaggCgggttctcacaccCtccagatgaigtatggctgcga  
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gaggacctgcgctcttggaccgcgccggacatggcGgctcagatcaccaagcgcaagtgaggaggcgcccatgtgg  
cggagcagcAgagagCctacctggagggcaCgtgcgtggacgggctccgcagatacctggagaacgggaaggagac  
gctgcagcgacgg(SEQ ID NO:121) ;

A\*2417 :

5 gctcccatccatgaggtatttctccacatccgtgtcccgcccgcccgcggggagccccgcttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagccagaggatggagccgcgggcgccgtgg  
atagagcaggaggggcccggagtattgggacgaggagacaGggaaagtgaaggcccatcacagactgaccgagaga  
acctgcggatcgcgctcCgtactacaaccagagcgaggCcggttctcacacctccagatgaigtTggctgcga  
cgtggggTcggacggcgcttccctccgcggtlaccggcaggacgcctacgacggcaaggattacatcgccctgaaA  
10 gaggacctgcgctcttggaccgcgccggacatggcGgctcagatcaccaagcgcaagtgaggaggcgcccatgtgg  
cggagcagcAgagagCctacctggagggcaCgtgcgtggacgggctccgcagatacctggagaacgggaaggagac  
gctgcagcgacgg(SEQ ID NO:122) ;

A\*2418 :

gctcccatccatgaggtatttctccacatccgtgtcccgcccgcccgcggggagccccgcttcatcgccgtggg  
15 ctacgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagccagaggatggagccgcgggcgccgtgg  
atagagcaggaggggcccggagtattgggacgaggagacagggaaagtgaaggcccatcacagactgaccgagaga  
acctgcggatcgcgctccgtactacaaccagagcgaggccggttctcacacctccagatgaigtTggctgcga  
cgtggggTcggacggcgcttccctccgcggtlaccaccagTaccgctacgacggcaaggattacatcgccctgaaA  
gaggacctgcgctcttggaccgcgccggacatggcggtcagatcaccaagcgcaagtgaggaggcgcccatgAgg  
20 cggagcagTTgagagcctacctggaTggcacgtgcgtggagtggtccgcagatacctggagaacgggaaggagac  
gctgcagcgacgg(SEQ ID NO:123) ;

A\*2419 :

gctcccatccatgaggtatttctccacatccgtgtcccgcccgcccgcggggagccccgcttcatcgccgtggg  
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25 atagagcaggaggggcccggagtattgggacgaggagacaGggaaagtgaaggcccaGtcacagactgaccgagtG  
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gaggacctgcgctcttggaccgcgccggacatggcGgctcagatcaccaagcgcaagtgggaggcggcccaigtgg  
cggagcagcAgagagCctaccitggagggcaCgtgcgtggacgggtccgcagataccitggagaacgggaaggagac  
gctgcagcgacgg(SEQ ID NO:124) ;

A\*2420 :

5 gctcccaatccatgaggtatttctccacatccgtgtcccgcccgccgaggagccccgcttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagccagaggatggagccgcgggcgccgtgg  
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acctgcggatcgcgctcGcttactacaaccagagcgaggCcggttctcacacccctccagatgatgtTggctgcga  
cgtggggTcggacggcgcttccctccgcggtlaccaccagTaccgctacgacggcaaggattacatcgccctgaaA  
10 gaggacctgcgctcttggaccgcgccggacatggcGgctcagatcaccaagcgcaagtgggaggcggcccaigtgg  
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gctgcagcgacgg(SEQ ID NO:125) ;

A\*2421 :

gctcccatccatgaggtatttctccacatccgtgtcccgcccgccgaggagccccgcttcatcgccgtggg  
15 ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagccagaggatggagccgcgggcgccgtgg  
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gaggacctgcgctcttggaccgcgccggacatggcGgctcagatcaccaagcgcaagtgggaggcggcccaigtgg  
20 cggagcagcAgagagCctaccitggagggcaCgtgcgtggacgggtccgcagataccitggagaacgggaaggagac  
gctgcagcgacgg(SEQ ID NO:126) ;

A\*2422 :

atggccgtcatggcggccgaacctcgttccgtactctcgggggcccgtggccctgaccagacctgggcaggct  
cccactccatgaggtatttctccacatccgtgtcccgcccgccgaggagccccgcttcatcgccgtgggcta  
25 cgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagccagaggatggagccgcgggcgccgtggata  
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gacctgcgctcttggaccgcggcggacatggcggctcagatcaccaagcgcaagtgaggagcggcccatgtggcgg  
agcagTggagagcttacctggagggcacgtgcgtggagTggctccgcagatacctggagaacgggaaggagacgct  
gcagcgcacggaccccccaagacacataatgaccaccaccccatctctgaccatgaggccacictgagatgctgg  
5 gcccitgggcttctaccctgcggagatcacactgacctggcagcgggaaggaggaccagaccaggacacggagc  
tttgggagaccaggccitgcagggatggaaccttccagaagtgggcAgctgtgggtggacaccttggagaggagca  
gagatacacctgccatgtgcagcatgagggctgcccaggccctcaccttgagatgggag (SEQ ID

NO:127) ;

A\*2423 :

10 gctccactccatgaggtatttctccacatccgtgtcccgcccgccgcgggagccccgcttcatcgccgtggg  
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atagagcaggaggggcccggagtattgggacgaggagacaGggaaagtgaaggcccacacacagatgaccgagaga  
acctgcggatcgcgctccgctactacaaccagagcgaggccggttctcacacctccagatgaltgttggctgcga  
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15 gaggacctgcgctcttggaccgcggcggacatggcggctcagatcaccaagcgcaagtgaggagcggcccatgtgg  
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gctgcagcgacgg (SEQ ID NO:128) ;

A\*2424 :

gctccactccatgaggtatttctccacatccgtgtcccgcccgccgcgggagccccgcttcatcgccgtggg  
20 ctacgtggacgacacgcagttcgtgcgggtcgacagcgacgccgcgagccagaggatggagcccgggcgccgtgg  
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acctgcggatcgcgctccgctactacaaccagagcgaggccggttctcacacctccagatgaltgtTggctgcga  
cgtggggctggacgggcgcttccctccggggtaccaccagtlacgcctacgacggcaaggattacatcgccctgaaA  
gaggacctgcgctcttggaccgcggcggacatggcggctcagatcacccagcgcaagtgaggagcggcccGtgTgg  
25 cggagcagTgagagCctacctggagggcaCgtgcgtggacgggtccgcagatacctggagaacgggaaggagac  
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A\*2425 :

gctcccatccatgaggtGtttctccacatccgtgtcccgcccgccgaggagccccgcttcatcgccgtggg  
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atagagcaggagggccggagtattgggacgaggagacaggaaagtgaagggccactcacagactgaccgagaga  
acctgcggatcgcgctccgtactacaaccagagcgaggccggtttcacacctccagatgatgtttggctgcga  
5 cgtggggtcggacggcgcttcciccgcggttaccaccagttacgcctacgacggcaaggattacatcgccctgaaa  
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gctgcagcgacgg(SEQ ID NO:130) ;

A\*2426 :

10 aacctctctctgctactctcgggggccctggccctgaccagacctgggcaggctcccatccatgaggtatitc  
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tgcggttcgacagcgacgcccgagccagaggatggagccgccccgcccgtggatagagcaggagggccggagta  
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tacaaccagagcgaggccggtttcacacctccagatgatgtttggctgcgacgtgggttcggacggcgcttcc  
15 tccAcgggttaccaccagttacgcctacgacggcaaggattacatcgccctgaaaggacctgcgctcttggaccgc  
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gagggcacgtgcgtggacgggtccgcagatacctggagaacgggaaggagacgctgcagcgacgg (SEQ ID  
NO:131) ;

A\*2427 :

20 atggccgtcatggcgccccgaacctctgtcttctctcgggggccctggccctgaccagacctgggcaggct  
cccatccatgaggtatitctccacatccgtgtcccgcccgcccgaggagccccgcttcatcgccgtgggtta  
cgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagccagaggatggagccgccccgcccgtggata  
gagcaggagggccggagtattgggacgaggagacaggaaagtgaagggccactcacagactgaccgagagaacc  
tgcggatcgcgctccgtactacaaccagagcgaggccggtttcacacctccagatgatgtttggctgcgacgt  
25 ggggtcggacggcgcttcciccgcggttaccaccagttacgcctacgacggcaaggattacatcgccctgaaagag  
gacctgcgctcttggaccgccccggacaGggcggtcagatcaccaagcgcaagtgggaggcggcccatgtggcgg  
agcagcagagagccctacctggagggcacgtgcgtggacgggtccgcagatacctggagaacgggaaggagacgct



gcagcgcacgg (SEQ ID NO:132) ;

A\*2428 :

gctccactccatgaggtatttctccacatccgtgtcccgcccgccgcggggagccccgcttcatcgccgtggg  
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5 atagagcaggagggggccggagtattgggacgaggagacaGggaaagigaaggcccactcacagactcaccgagtG  
acctggggacccitcgcggttactacaaccagagcgaggCcggttctcacacctccagatgaigtTggctgcga  
cgtggggTcggacgggcgcttctcccggggtaccaccagTaccctacgacggcaaggattacatgcccigaaA  
gaggacctgcgtcttggaccgcggcgacatggcGgctcagatcaccaagcgcaagtgggaggcgcccatgttg  
cggagcagcAgagagCctacctggaggggcaCgtgcgtggacgggtccgcagatacctggagaacgggaaggagac  
10 gctgcagcgcacgg (SEQ ID NO:133) ;

A\*2429 :

gctccactccatgaggtatttctccacatccgtgtcccgcccgccgcggggagccccgcttcatcgccgtggg  
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15 acctgcggatcgcgtcCgctactacaaccagagcgaggCcggttctcacacctccagatgaigtTggctgcga  
cgtggggTcggacgggcgcttctcccggggtaccaccagTaccctacgacggcaaggattacatgcccigaaA  
gaggacctgcgtcttggaccgcggcgacatggcGgctcagatcaccaagcgcaagtgggaggcgcccatgttg  
cggagcagcAgagagCctacctggaggggcaCgtgcgtggacgggtccgcagatacctggagaacgggaaggagac  
gctgcagcgcacgg (SEQ ID NO:134) ;

20 A\*2430 :

gctccactccatgaggtatttctccacatccgtgtcccgcccgccgcggggagccccgcttcatcgccgtggg  
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atagagcaggagggggccggagtattgggacgaggagacaGggaaagigaaggcccactcacagactCaccgagaga  
acctgcggatcgcgtcCgctactacaaccagagcgaggCcggttctcacacctccagatgaigtTggctgcga  
25 cgtggggTcggacgggcgcttctcccggggtaccaccagTaccctacgacggcaaggattacatgcccigaaA  
gaggacctgcgtcttggaccgcggcgacatggcGgctcagatcaccaagcgcaagtgggaggcgcccatgttg  
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gctgcagcgacgg(SEQ ID NO:135) ;

A\*2431 :

gctcccatccatgaggtatttctccacatccgtgtcccgcccgccgaggagccccgcttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagccagaggatggagccgaggcgccgtgg  
5 atagagcaggaggggcccggagiatgggacgagCagacagggaaggaaggccctcacagactgaccgagaga  
acctgcggatcgcgtccgctactacaaccagagcgaggccggtttcacacctccagatgatgttggctgcga  
cgtggggctggacggcgcttccctccgcggtaccaccagtagccctacgacggcaaggattacatcgccctgaaa  
gaggacctgcgtcttggaccgcgccggacatggcggctcagatcaccaagcgcaagtgggaggcgcccaatTgg  
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10 gctgcagcgacgg(SEQ ID NO:136) ;

A\*2432 :

gctcccatccatgaggtatttctccacatccgtgtcccgcccgccgaggagccccgcttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagccagaggatggagccgaggcgccgtgg  
atagagcaggaggggcccggagiatgggacgaggagacagggaaggaaggccctcacagactgaccgagaga  
15 Gcctgcggatcgcgtccgctactacaaccagagcgaggccggtttcacacctccagatgatgttggctgcga  
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gaggacctgcgtcttggaccgcgccggacatggcggctcagatcaccaagcgcaagtgggaggcgcccaatTgg  
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20 A\*2433 :

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A\*2434 :

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15 A\*2435 :

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A\*2438 :

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15 A\*2501 :

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25 A\*2504 :

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A\*2601 :

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A\*2602 :

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A\*2603 :

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NO:149) ;

A\*2604 :

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25 cgtggacgacacgcagttcgttgcggttcgacagcgacgccgcgagccagaggatggagccgcggcgccgtggata  
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A\*2605 :

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NO:151) ;

A\*2606 :

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5 A\*2607 :

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A\*2608 :

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A\*2609 :  
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15 A\*2610 :  
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A\*2613 :

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A\*2615 :

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A\*2904 :

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A\*2905 :

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A\*2906 :

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A\*2907 :

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A\*3001 :

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NO:171) ;

A\*3002 :

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NO:172) ;

A\*3003 :

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A\*3004 :

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A\*3006 :

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A\*3007 :

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A\*3008 :

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A\*3009 :  
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A\*3010 :  
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A\*3011 :

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A\*3012 :

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25 A\*310102 :

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A\*3102 :  
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A\*3103 :  
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A\*3104 :

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A\*3105 :

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A\*3106 :

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A\*3107 :

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A\*3108 :

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A\*3109 :

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A\*3201 :

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A\*3204 :
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A\*3205 :

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A\*3206 :

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A\*3207 :

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5 A\*3301 :

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A\*3303 :

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A\*3304 :

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15 A\*3305 :

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A\*3306 :

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A\*3401 :

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A\*3402 :

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A\*3403 :

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A\*3404 :

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A\*3405 :

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A\*3601 :

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A\*3602 :

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5 A\*3603 :

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A\*3604 :

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A\*4301 :

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15 A\*6601 :

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25 A\*6604 :

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A\*680101 :

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gcagcgcacgg (SEQ ID NO:243) ;

5 A\*7404 :

ggctcccactccaatgaggtatcttccacatccgtgtcccgccccggcggggagccccgttcatcgccgtgg  
gctacgtggacgacacgcagttcgtgcggtttgacagcgacgccgagccagaggatggagccgcgggcgccgtg  
gatagagcaggaggggtccggagtattgggacggggagacacggaaAgtgaaggccctcacagactgaccgagt  
Gacctggggacccctgcgcggctactacaaccagagcgaggccggttctcacacatccagatgaigtatggctgcg  
10 acgtggggccggacgggcgcCtctccgcggtaccagcaggacgcctacgacggcaaggattacatcgccctgaa  
cgaggacctgcgctcttggaccgcgggcgacatggcggtcagatcaccagcgcaagtgggaggcgcccggtt  
gcggagcagttgagagcctacctggagggcacgtgctggagtggctccgcagatacctggagaacgggaaggaga  
cgctgcagcgcacgg (SEQ ID NO:244) ;

A\*7405 :

15 gctcccactccaatgaggtatcttccacatccgtgtcccgccccggcggggagccccgttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggtttgacagcgacgccgagccagaggatggagccgcgggcgccgtg  
atagagcaggagggggccggagtattgggaccaggagacacggaatgtgaaggccctcacagGctgaccgagtgg  
acctggggacccctgcgcggctactacaaccagagcgaggccggttctcacacatccagatgaigtatggctgcga  
cggtggggccggacgggcgcctcctccgcggtaccagcaggacgcctacgacggcaaggattacatcgccctgaa  
20 gaggacctgcgctcttggaccgcgggcgacatggcggtcagatcaccagcgcaagtgggaggcgcccggttgg  
cggagcagttgagagcctacctggagggcacgtgctggagtggctccgcagatacctggagaacgggaaggagac  
gctgcagcgcacgg (SEQ ID NO:245) ;

A\*7406 :

25 gctcccactccaatgaggtatcttccacatccgtgtcccgccccggcggggagccccgttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggtttgacagcgacgccgagccagaggatggagccgcgggcgccgtg  
atagagcaggagggggccggagtattgggaccaggagacacggaatgtgaaggccctcacagactCaccgagtG  
acctggggacccctgcgcggctactacaaccagagcgaggccggttctcacacatccagatgaigtatggctgcga

cgtggggccggacggggcgCtctccgcggtaccagcaggacgcctacgacggcaaggattacatcgccitgaac  
gaggaccctgcgcctctggaccgcgggcgacatggcggctcagatcaccagcgcaagtgggaggcggcccGtgtgg  
cggagcagTtagagcctacctggagggcacgtgcgtggagtggciccgcagatactggagaacgggaaggagac  
gctgcagcgacg(SEQ ID NO:246) ;

5 A\*7407 :

gctcccactccaatgaggtatttcttcacatccgtgtcccggcccgccgaggagccccgttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggtttgacagcgacgccgcgagccagaggatggagccgcgggcgccgtgg  
atagagcaggagggggccggagtattgggaccaggagacacggaatgtgaagggccactcacagaTtgaccgagtG  
acctggggacccctgcgcggctactacaaccagagcgaggccggttctcacacatccagatgatgtatggctgcga  
10 cgtggggccggacggggcgCtctccgcggtaccagcaggacgcctacgacggcaaggattacatcgccitgaac  
gaggaccctgcgcctctggaccgcgggcgacatggcggctcagatcaccagcgcaagtgggaggcggcccGtgtgg  
cggagcagTtagagcctacctggagggcacgtgcgtggagtggciccgcagatactggagaacgggaaggagac  
gctgcagcgacgg(SEQ ID NO:247) ;

A\*7408 :

15 gctcccactccaatgaggtatttcttcacatccgtgtcccggcccgccgaggagccccgttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggtttgacagcgacgccgcgagccagaggatggagccgcgggcgccgtgg  
atagagcaggagggggccggagtattgggaccaggagacacggaatgtgaagggccactcacagactgaccgagtgg  
acctggggacccctgcgcggctactacaaccagagcgaggccggttctcacacatccagatgatgtatggctgcga  
cgtggggccggacggggcgCtctccgcggtaccagcaggacgcctacgacggcaaggattacatcgccitgaac  
20 gaggaccctgcgcctctggaccgcgggcgacatggcggctcagatcaccagcgcaagtgggaggcggccAgtgtgg  
cggagcagTtagagcctacctggagggcacgtgcgtggagtggciccgcagatactggagaacgggaaggagac  
gctgcagcgacgg(SEQ ID NO:248) ;

A\*7409 :

25 gctcccactccaatgaggtatttcttcacatccgtgtcccGccccggcccggggagccccgttcatcgccgtggg  
ctacgtggacgacacgcagttcgtgcggtttgacagcgacgccgcgagccagaggatggagccgcgggcgccgtgg  
atagagcaggagggggccggagtattgggaccaggagacacggaatgtgaagggccactcacagactgaccgagtgg  
acctggggacccctgcgcggctactacaaccagagcgaggccggttctcacacatccagatgatgtatggctgcga

cgtggggccggacgggcgccctcccggggtaccagcaggacgcciacgacggcaaggattacatcgccitgaac  
 gaggacctgcgtcttggaccgcgccggacatggcggctcagatcacccagcgcaagtgggaggcgcccggtgg  
 cggagcagttgagagcctacctggaggcgacgtgcgtggagtggctccgcagatacctggagaacgggaaggagac  
 gctgcagcgacgg (SEQ ID NO:249) ;

5 A\*8001 :

Atggccgtcatgcccccgaacctccctcctgtctactctcgggggccctggccctgaccagacctgggcaggct  
 cccactccatgaggtatctctcacatccgtgtccggcccgccggggagccccgcctcatcgcatgggcta  
 cgtggacgactcgcagttcgtgcagttcgacagcgacgccgagccagaggatggagccgcgggcgccgtggata  
 gagcaggaggagccggagtattgggacgaggagacacggaatgtgaaggcccactcacagactaaccgagcgaacc  
 10 tggggacctgcgcggctactacaaccagagcgaggacggcttctcacaccatccagataatgtatggctgcgacgt  
 ggggtcggacggcgcttccctccgcggtaccggcaggacgcttacgacggcaaggattacatcgccctgaacgag  
 gacctgcgtcttggaccgcgccggacatggcggctcagatcaccaagcgcaagtgggaggcgcccgctgggcgg  
 agcagctgagagcctacctggaggcgagtgctggacgggctccgcagatacctggagaacgggaaggagacgt  
 gcagcgacggaccccccaagacacataigaccaccaccccatctctgacctgaggccactctgaggtgctgg  
 15 gcccgtagcttctacctgcggagatcacactgacctggcagcgggatggggaggaccagaccaggacacggagc  
 tctggagaccaggccctgcaggggatggaaccttccagaagtggcgccgtgtgggtggtaaccttctggaaaggagaa  
 gagataacctgccatgtgcagcatgagggtctgcccGagccctcacctgagatgggag  
 (SEQ ID NO:25) ;

20

The probe list A1 is shown in Tables 1-1 to 1-7  
 and the probe list A2 is shown in Tables 2-1 to 2-6.  
 The allele-probe lists are shown in Tables 3-1 to 3-9  
 and Tables 4-1 to 4-9.

25



Table 1-1

Probe No.	Base Sequence
0	g ccc cgc ttc atc gcC (SEQ ID No: 2 5 1)
1	gac cag gag aca cgg aat A (SEQ ID No: 2 5 2)
2	gcg gag cag cgg aga gT (SEQ ID No: 2 5 3)
3	a gtc tac ctg gag ggc C (SEQ ID No: 2 5 4)
4	gtc tac ctg gag ggc cG (SEQ ID No: 2 5 5)
5	agg tgc tgg gcc ctg G (SEQ ID No: 2 5 6)
6	g gtg gtg cct tct gga G (SEQ ID No: 2 5 7)
7	c acc ctg aga tgg gag cT (SEQ ID No: 2 5 8)
8	cc ctg aga tgg gag ctG (SEQ ID No: 2 5 9)
9	g gac atg gca gct cag atT (SEQ ID No: 2 6 0)
10	cac tcc atg agg tat ttc tC (SEQ ID No: 2 6 1)
11	c cgg ccc ggc agt ggA (SEQ ID No: 2 6 2)
12	t tct cac acc atc cag atG (SEQ ID No: 2 6 3)
13	c cat gcg gcg gag cag T (SEQ ID No: 2 6 4)
14	cat gcg gcg gag cag tT (SEQ ID No: 2 6 5)
15	ata gag cag gag agg ccT (SEQ ID No: 2 6 6)
16	c tca cag act gac cga gA (SEQ ID No: 2 6 7)
17	c tac aac cag agc gag gC (SEQ ID No: 2 6 8)
18	ga gtc tac ctg gag ggc T (SEQ ID No: 2 6 9)
19	gtg gac gac acg cag ttA (SEQ ID No: 2 7 0)
20	tg cta ctc tcg ggg gcT (SEQ ID No: 2 7 1)
21	g gcc cac tca cag act C (SEQ ID No: 2 7 2)
22	g gcc ggt tct cac acc G (SEQ ID No: 2 7 3)
23	t tct cac acc gtc cag aG (SEQ ID No: 2 7 4)
24	c gac gtg ggg tcg gac T (SEQ ID No: 2 7 5)
25	gg gag gcg gcc cat gT (SEQ ID No: 2 7 6)
26	c cat gtg gcg gag cag tT (SEQ ID No: 2 7 7)
27	gcc tac ctg gag ggc aC (SEQ ID No: 2 7 8)
28	ga gct gtg gtc gct gcT (SEQ ID No: 2 7 9)
29	ag ccc cgc ttc atc gcA (SEQ ID No: 2 8 0)
30	ccg gag tat tgg gac gG (SEQ ID No: 2 8 1)

Table 1-2

Probe No.	Base Sequence
31	ggc ttg cat tcc ctc cG (SEQ ID No :32)
32	c cca gtt ggg acg agt gT (SEQ ID No :33)
33	ct gct gct gct gct gcT (SEQ ID No :34)
34	a gaa gat gtc ctg gga aaC (SEQ ID No :35)
35	t gtg cag tca ggg ttt ctT (SEQ ID No :36)
36	gcc tca gag ggc aac atC (SEQ ID No :37)
37	ct gct gct gct gct gcT (SEQ ID No :38)
38	ttc tat ccc cgg aat atc aT (SEQ ID No :39)
39	gtt gct gct gct gct gcT (SEQ ID No :40)
40	cag acc ttg gcc atg aac A (SEQ ID No :41)
41	gg aat cac agc act cac G (SEQ ID No :42)
42	a cgg cga tat cta aaa tcc A (SEQ ID No :43)
43	ctc tcc caa aac ctg gag T (SEQ ID No :44)
44	ttc ttg aag gaa gat gcc G (SEQ ID No :45)
45	cat gaa gac aac agc acc aA (SEQ ID No :46)
46	ggg ttt ctc gct gag gG (SEQ ID No :47)
47	caa gga gag gag cag agT (SEQ ID No :48)
48	g gcc acc agg att tgc G (SEQ ID No :49)
49	c agg gct tct ggc ttc tG (SEQ ID No :50)
50	ag aaa aca tca gct gca gaT (SEQ ID No :51)
51	at caa cac cca gtt ggg aT (SEQ ID No :52)

Table 1-3

Probe No.	Base Sequence
61	ca cag act cac cga gtg G ( SEQ ID No: 3 1 2 )
62	c gcg gcg gac atg gcG ( SEQ ID No: 3 1 3 )
63	gt ccg gag tat tgg gac G ( SEQ ID No: 3 1 4 )
64	ac ggg gag aca cgg aaC ( SEQ ID No: 3 1 5 )
65	ca gtg ggc tac gtg gac A ( SEQ ID No: 3 1 6 )
66	tgg gag acg gcc cat gT ( SEQ ID No: 3 1 7 )
67	c cat gag gcg gag cag tT ( SEQ ID No: 3 1 8 )
68	a gct cag acc acc aag cA ( SEQ ID No: 3 1 9 )
69	cat gcg gcg gag cag cA ( SEQ ID No: 3 2 0 )
70	cg tgg ata gag cag gag A ( SEQ ID No: 3 2 1 )
71	gac ggg gag aca cgg C ( SEQ ID No: 3 2 2 )
72	c tgg gcg ggc tct caG ( SEQ ID No: 3 2 3 )
73	tc gac agc gac gcc gG ( SEQ ID No: 3 2 4 )
74	c acc gtc cag agg atg tC ( SEQ ID No: 3 2 5 )
75	cgg aaa gtg aag gcc caG ( SEQ ID No: 3 2 6 )
76	g gcc cag tca cag act C ( SEQ ID No: 3 2 7 )
77	g gct cag atc acc aag cA ( SEQ ID No: 3 2 8 )
78	gcg gag cag ttg aga gC ( SEQ ID No: 3 2 9 )
79	g ggc acg tgc gtg gaG ( SEQ ID No: 3 3 0 )
80	g tgg gag gcg gcc cG ( SEQ ID No: 3 3 1 )
81	gg gag gcg gcc cgt gT ( SEQ ID No: 3 3 2 )
82	c cgc ggg tac cag cag T ( SEQ ID No: 3 3 3 )
83	g gag ccc cgc ttc atc T ( SEQ ID No: 3 3 4 )
84	gac cag gag aca cgg aaA ( SEQ ID No: 3 3 5 )
85	at tgg gac gag gag aca G ( SEQ ID No: 3 3 6 )
86	gac gag gag aca ggg aaA ( SEQ ID No: 3 3 7 )
87	g aag gcc cac tca cag aG ( SEQ ID No: 3 3 8 )
88	g agg tat ttc ttc aca tcc A ( SEQ ID No: 3 3 9 )
89	ttc ctc cgc ggg tat gaA ( SEQ ID No: 3 4 0 )
90	gag tat tgg gac cgg aaC ( SEQ ID No: 3 4 1 )

Table 1-4

Probe No.	Base Sequence
91	cgg aat gtg aag gcc caG (SEQ ID No: 3 4 2)
92	g gcc ggt tct cac acc C (SEQ ID No: 3 4 3)
93	t tct cac acc ctc cag aG (SEQ ID No: 3 4 4)
94	c cgg ccc ggc cgc gA (SEQ ID No: 3 4 5)
95	cgc ggg tac cac cag tT (SEQ ID No: 3 4 6)
96	ca cag act gac cga gtg G (SEQ ID No: 3 4 7)
97	g ttg aga gcc tac ctg gaT (SEQ ID No: 3 4 8)
98	cat gag gcg gag cag cT (SEQ ID No: 3 4 9)
99	ctg aga gcc tac ctg gaT (SEQ ID No: 3 5 0)
100	tgg ata gag cag gag ggT (SEQ ID No: 3 5 1)
101	cag aga gcc tac ctg gaT (SEQ ID No: 3 5 2)
102	ggc ctg gtt ctc ctt gC (SEQ ID No: 3 5 3)
103	g aga gcc tac ctg gat gC (SEQ ID No: 3 5 4)
104	ggc tgc gac gtg ggg T (SEQ ID No: 3 5 5)
105	g ggc cgg tgc gtg gaG (SEQ ID No: 3 5 6)
106	ggc cgg tgc gtg gag T (SEQ ID No: 3 5 7)
107	gc tct tgg acc gcg gcA (SEQ ID No: 3 5 8)
108	gg ccc ggc cgc ggg A (SEQ ID No: 3 5 9)
109	gg gag gcg gcc cgt gA (SEQ ID No: 3 6 0)
110	cgt gag gcg gag cag cA (SEQ ID No: 3 6 1)
111	g gca gct cag atc acc G (SEQ ID No: 3 6 2)
112	g ccg gac ggg cgc ttA (SEQ ID No: 3 6 3)
113	g cag aga gcc tac ctg C (SEQ ID No: 3 6 4)
114	g ccg gag tat tgg gac cT (SEQ ID No: 3 6 5)
115	g gca gct cag atc acc aG (SEQ ID No: 3 6 6)
116	g gag gcg gcc cgt cG (SEQ ID No: 3 6 7)
117	ac gag gag aca ggg aaa G (SEQ ID No: 3 6 8)
118	cc cag ccc acc gtc cA (SEQ ID No: 3 6 9)
119	c cgt gtg gcg gag cag T (SEQ ID No: 3 7 0)
120	gcg gag cag tgg aga gC (SEQ ID No: 3 7 1)

Table 1-5

Probe No.	Base Sequence
121	ggc aag gat tac atc gcc T (SEQ ID No: 372)
122	cgt gtg gcg gag cag tT (SEQ ID No: 373)
123	c tcc cac tcc atg agg tG (SEQ ID No: 374)
124	cg ctc cgc tac tac aac G (SEQ ID No: 375)
125	ctg cgg atc gcg ctc C (SEQ ID No: 376)
126	gcg gag cag cag aga gC (SEQ ID No: 377)
127	a tct tcc cag ccc acc G (SEQ ID No: 378)
128	ctg ggc ttc tac cct gcA (SEQ ID No: 379)
129	cgc ggg tac cac cag taT (SEQ ID No: 380)
130	ag acg ctg cag cgc acT (SEQ ID No: 381)
131	g gcg gct cag atc acc C (SEQ ID No: 382)
132	ggg aaa gtg aag gcc caG (SEQ ID No: 383)
133	cc tgg gca ggc tcc caA (SEQ ID No: 384)
134	g ggc acg tgc gtg gac T (SEQ ID No: 385)
135	gac ggg cgc ttc ctc cA (SEQ ID No: 386)
136	gg acc gcg gcg gac aG (SEQ ID No: 387)
137	cg gag tat tgg gac gag C (SEQ ID No: 388)
138	a cag act gac cga gag aG (SEQ ID No: 389)
139	c cag agg atg gag ccg T (SEQ ID No: 390)
140	g agc cag agg atg gag cT (SEQ ID No: 391)
141	gc tcc cac tcc atg agC (SEQ ID No: 392)
142	g cct gca ggg gat ggG (SEQ ID No: 393)
143	c cag cgc aag tgg gag A (SEQ ID No: 394)
144	c cgc ggg tac cag cag A (SEQ ID No: 395)
145	gcc tac ctg gag ggc cT (SEQ ID No: 396)
146	tc cgc ggg tac cag cG (SEQ ID No: 397)
147	ttc ctc cgc ggg tac cA (SEQ ID No: 398)
148	gg tac cag cag gac gcT (SEQ ID No: 399)
149	cg cag ttc gtg cgg ttG (SEQ ID No: 400)
150	c cag agc gag gac ggt A (SEQ ID No: 401)

Table 1-6

Probe No.	Base Sequence
151	cag atg atg tat ggc tgc C ( SEQ ID No: 4 0 2 )
152	g atg gag ccg cgg gcA ( SEQ ID No: 4 0 3 )
153	g gac ctg cag aca cgg C ( SEQ ID No: 4 0 4 )
154	gag acg ctg cag cgc G ( SEQ ID No: 4 0 5 )
155	tgg gag gcg gcc cgt T ( SEQ ID No: 4 0 6 )
156	gg gag gcg gcc cgt C ( SEQ ID No: 4 0 7 )
157	g ggc tac gtg gac gac G ( SEQ ID No: 4 0 8 )
158	cac acc atc cag ata atg C ( SEQ ID No: 4 0 9 )
159	gtg cag cat gag ggt ctC ( SEQ ID No: 4 1 0 )
160	gg tac ccg cag gac gcT ( SEQ ID No: 4 1 1 )
161	c cac tcc atg agg tat ttc A ( SEQ ID No: 4 1 2 )
162	g aca ccg aat gtg aag gG ( SEQ ID No: 4 1 3 )
163	c cta gtt ctc itt gga gct A ( SEQ ID No: 4 1 4 )
164	gg ccg gac ggg cgc C ( SEQ ID No: 4 1 5 )
165	gcc tac ctg gat ggc aC ( SEQ ID No: 4 1 6 )
166	t ggc acg tgc gtg gag T ( SEQ ID No: 4 1 7 )
167	gac cag gag aca ggg aaA ( SEQ ID No: 4 1 8 )
168	gc acg gac ccc ccc aG ( SEQ ID No: 4 1 9 )
169	ac gag gac ctg agc tcC ( SEQ ID No: 4 2 0 )
170	gcg ccg tgg ata gag cG ( SEQ ID No: 4 2 1 )
171	g ccg gcg ccg tgg atG ( SEQ ID No: 4 2 2 )
172	c ccc atc gtg ggc atc C ( SEQ ID No: 4 2 3 )
173	ctg cag cgc acg gac G ( SEQ ID No: 4 2 4 )
174	g gac gcc ccc aag acG ( SEQ ID No: 4 2 5 )
175	ctc ttt gga gct gtg atc G ( SEQ ID No: 4 2 6 )
176	gac ggc aag gat tac atc T ( SEQ ID No: 4 2 7 )
177	gtc tac ctg gag ggc aC ( SEQ ID No: 4 2 8 )
178	cgg aga gcc tac ctg gaT ( SEQ ID No: 4 2 9 )
179	g gac ggt tct cac acc C ( SEQ ID No: 4 3 0 )
180	g ggc gag tgc gtg gag T ( SEQ ID No: 4 3 1 )

Table 1-7

Probe No.	Base Sequence
181	g gag tgg ctc cgc aga C (SEQ ID No.:4 3 2)
182	ga acc ttc cag aag tgg gT (SEQ ID No.:4 3 3)
183	cc atg agg tat ttc tac acT (SEQ ID No.:4 3 4)
184	g agg tat ttc tac acc tcc A (SEQ ID No.:4 3 5)
185	cgc ggg tac cgg cag C (SEQ ID No.:4 3 6)
186	cat gtg gcg gag cag cT (SEQ ID No.:4 3 7)
187	g ccg gag tat tgg gac G (SEQ ID No.:4 3 8)
188	ag tgg gag gcg gcc cT (SEQ ID No.:4 3 9)
189	gc ggg tac cgg cag gT (SEQ ID No.:4 4 0)
190	tgg aga gcc tac ctg gaT (SEQ ID No.:4 4 1)
191	tg ggg tcg gac ggg cA (SEQ ID No.:4 4 2)
192	gc aga tac ctg gag aac C (SEQ ID No.:4 4 3)
193	gac ctg ggg acc ctg cA (SEQ ID No.:4 4 4)
194	gt tct cac acc atc cag aG (SEQ ID No.:4 4 5)
195	g gcc ctg acc cag acc A (SEQ ID No.:4 4 6)
196	c ctc ctc ctg cta ctc tT (SEQ ID No.:4 4 7)
197	ctc ctc cgc ggg tac cA (SEQ ID No.:4 4 8)
198	gac cga gtg gac ctg gC (SEQ ID No.:4 4 9)
199	g aag gcc cac tca cag G (SEQ ID No.:4 5 0)
200	ca cag att gac cga gtg G (SEQ ID No.:4 5 1)
201	c aag tgg gag gcg gcc A (SEQ ID No.:4 5 2)
202	c ttc aca tcc gtg tcc cC (SEQ ID No.:4 5 3)
203	cag ccc acc atc ccc atT (SEQ ID No.:4 5 4)

Table 2-1

Probe No.	Base Sequence
0	a gag acc agA gac ttg aca (SEQ ID No:53)
1	ctg gag act Aag gaa tgg a (SEQ ID No:54)
2	cga tat cta Aaa tcc ggc g (SEQ ID No:55)
3	cta aaa tcc Ggc gta gtc c (SEQ ID No:56)
4	c aca ctg aGc tgg cgt c (SEQ ID No:57)
5	att att ttc taC gtc tgt tgt t (SEQ ID No:58)
6	tg ctg tcc Ggg gat gga (SEQ ID No:59)
7	acc cgc agT gag gcc tc (SEQ ID No:60)
8	g agg aga aGa gtg ccc c (SEQ ID No:61)
9	tg atg tca gCt ctt ggg tc (SEQ ID No:62)
10	c ctg cgc tAt gac agg c (SEQ ID No:63)
11	gaa tgg aca Gtg ccc cag (SEQ ID No:64)
12	c aca ctg aCc tgg cgt c (SEQ ID No:65)
13	gg att tgc cGa gga gag g (SEQ ID No:66)
14	gaa tcc agc Ata gtc ctg a (SEQ ID No:67)
15	a gag acc agG gac ttg ac (SEQ ID No:68)
16	ctg gag act Gag gaa tgg (SEQ ID No:69)
17	gtt gct gct G gct gct g (SEQ ID No:70)
18	g gtg gcc acT agg att tg (SEQ ID No:71)
19	gct gct g gct gct gcT a (SEQ ID No:72)
20	agc gag gcA tca gag gg (SEQ ID No:73)
21	tcc caa aac Gtg gag act g (SEQ ID No:74)
22	at ttc tac taT gat ggg gag (SEQ ID No:75)
23	cta gaa tcc Agc gta gtc c (SEQ ID No:76)
24	t ggg tcc Gct ggc tcc (SEQ ID No:77)
25	cc aag aca cTc tat cac gc (SEQ ID No:78)
26	a gag gag caA agg ttc acc (SEQ ID No:79)
27	cga tat cta Gaa tcc ggc g (SEQ ID No:80)
28	tac tac gat Agg gag ctc t (SEQ ID No:81)
29	g ggt cca gGg ctc gtg (SEQ ID No:82)
30	cag gat ggg Cta tct ttg a (SEQ ID No:83)



Table 2-2

Probe No.	Base Sequence
31	at tcc ctc cGg gag att ag (SEQ ID No: 84)
32	t gct gct gct gct gcT at (SEQ ID No: 85)
33	ct gct gct gcT att ttt gtt (SEQ ID No: 86)
34	c ctg gga aaC aag aca tgg (SEQ ID No: 87)
35	a ggg ttt ctT gct gag gta (SEQ ID No: 88)
36	g ggc aac atC acc gtg ac (SEQ ID No: 89)
37	gct gct gct gct gcT att (SEQ ID No: 90)
38	cgg aat atc aTa ctg acc tg (SEQ ID No: 91)
39	gcc atg aac Atc agg aat tt (SEQ ID No: 92)
40	gc act cac Gct gtg ccc (SEQ ID No: 93)
41	cta aaa tcc Agc gta gtc c (SEQ ID No: 94)
42	aac ctg gag Tct gag gaa t (SEQ ID No: 95)
43	gaa gat gcc Gtg aag acc (SEQ ID No: 96)
44	c agc acc aAg agc tcc c (SEQ ID No: 97)
45	c gct gag gGa cat ctg g (SEQ ID No: 98)
46	g gag cag agT ttc acc tg (SEQ ID No: 99)
47	agg att tgc Gaa gga gag g (SEQ ID No: 100)
48	ct ggc ttc tGt ccc tgg a (SEQ ID No: 101)
49	a gct gca gaT ggt cca ga (SEQ ID No: 102)
50	ca gtt ggg aTg agt gac c (SEQ ID No: 103)

Table 2-3

Probe No.	Base Sequence
61	g aca cgg aaC gtg aag gc ( SEQ ID No : 516 )
62	tac gtg gac Aac acg cag ( SEQ ID No : 517 )
63	cc acc aag cAc aag tgg g ( SEQ ID No : 518 )
64	ag cag gag Agt ccg gag ( SEQ ID No : 519 )
65	gag aca cgg Caa gtg aag ( SEQ ID No : 520 )
66	g ggc tct caG tcc atg ag ( SEQ ID No : 521 )
67	c gac gcc gGg agc cag ( SEQ ID No : 522 )
68	g agg atg tCt ggc tgc g ( SEQ ID No : 523 )
69	g aag gcc caG tca cag ac ( SEQ ID No : 524 )
70	tc acc aag cAc aag tgg g ( SEQ ID No : 525 )
71	ag ttg aga gCc tac ctg g ( SEQ ID No : 526 )
72	tgc gtg gaG tgg ctc cg ( SEQ ID No : 527 )
73	gcg gcc cGt gtg gcg ( SEQ ID No : 528 )
74	g gcc cgt gTg gcg gag ( SEQ ID No : 529 )
75	tac cag cag Tac gcc tac ( SEQ ID No : 530 )
76	cgc ttc atc Tca gtg ggc ( SEQ ID No : 531 )
77	gag gag aca Ggg aaa gtg ( SEQ ID No : 532 )
78	g aca ggg aaA gtg aag gc ( SEQ ID No : 533 )
79	ac tca cag aGt cac cga g ( SEQ ID No : 534 )
80	ttc aca tcc Atg tcc cgg ( SEQ ID No : 535 )
81	c ggg tat gaA cag cac gc ( SEQ ID No : 536 )
82	g gac cgg aaC aca cgg aa ( SEQ ID No : 537 )
83	tct cac acc Ctc cag atg ( SEQ ID No : 538 )
84	ct cac acc Ctc cag agg ( SEQ ID No : 539 )
85	cc ctc cag aGg atg tat g ( SEQ ID No : 540 )
86	ggc cgc gAg gag ccc ( SEQ ID No : 541 )
87	c cac cag tTc gcc tac g ( SEQ ID No : 542 )
88	c tac ctg gaT ggc acg tg ( SEQ ID No : 543 )
89	g gag cag cTg aga gcc t ( SEQ ID No : 544 )
90	cag gag ggT ccg gag ta ( SEQ ID No : 545 )

Table 2-4

Probe No.	Base Sequence
91	ctg gag aac Cgg aag gag ( SEQ ID No : 5 4 6 )
92	c ctg gat gCc acg tgc g ( SEQ ID No : 5 4 7 )
93	c gtg ggg Tcg gac ggg ( SEQ ID No : 5 4 8 )
94	acc gcg gcA gac atg gc ( SEQ ID No : 5 4 9 )
95	c cgc ggg Aag ccc cg ( SEQ ID No : 5 5 0 )
96	gcg gcc cGt gag gcg ( SEQ ID No : 5 5 1 )
97	g gcc cgt gAg gcg gag ( SEQ ID No : 5 5 2 )
98	cag atc acc Gag cgc aag ( SEQ ID No : 5 5 3 )
99	ggg cgc tTA ctc cgc g ( SEQ ID No : 5 5 4 )
100	c tac ctg Cag gcc cgg ( SEQ ID No : 5 5 5 )
101	at tgg gac cTg cag aca c ( SEQ ID No : 5 5 6 )
102	ag atc acc aGg cgc aag t ( SEQ ID No : 5 5 7 )
103	gcc cgt cGg gcg gag ( SEQ ID No : 5 5 8 )
104	aca ggg aaa Gtg aag gcc ( SEQ ID No : 5 5 9 )
105	g aag tgg gcA gct gtg gt ( SEQ ID No : 5 6 0 )
106	g tgg aga gCc tac ctg g ( SEQ ID No : 5 6 1 )
107	tac atc gcc Ttg aac gag g ( SEQ ID No : 5 6 2 )
108	cc atg agg tGt ttc tcc ac ( SEQ ID No : 5 6 3 )
109	tac tac aac Gag agc gag g ( SEQ ID No : 5 6 4 )
110	tc gcg ctc Cgc tac tac ( SEQ ID No : 5 6 5 )
111	g cag aga gCc tac ctg g ( SEQ ID No : 5 6 6 )
112	c tac cct gcA gag atc ac ( SEQ ID No : 5 6 7 )
113	c cac cag taT gcc tac ga ( SEQ ID No : 5 6 8 )
114	cag atc acc Cag cgc aag ( SEQ ID No : 5 6 9 )
115	a ggc tcc caA tcc atg ag ( SEQ ID No : 5 7 0 )
116	t gtg gtg gTA cct tct gg ( SEQ ID No : 5 7 1 )
117	cg gag cag Tgg aga gtc ( SEQ ID No : 5 7 2 )
118	c gtg gac Tgg ctc cgc ( SEQ ID No : 5 7 3 )
119	c ttc ctc cAc ggg tac c ( SEQ ID No : 5 7 4 )
120	g gcg gac aGg gcg gct ( SEQ ID No : 5 7 5 )

Table 2-5

Probe No.	Base Sequence
121	tca cag act Cac cga gag ( SEQ ID No : 5 7 6 )
122	gg gac gag Cag aca ggg ( SEQ ID No : 5 7 7 )
123	c cga gag aGc ctg cgg ( SEQ ID No : 5 7 8 )
124	ac tca cag aTt gac cga ga ( SEQ ID No : 5 7 9 )
125	g gag ccg Tgg gcg cc ( SEQ ID No : 5 8 0 )
126	g atg gag cTg cgg gcg ( SEQ ID No : 5 8 1 )
127	c tcc atg agC tat ttc tcc ( SEQ ID No : 5 8 2 )
128	ggg gat ggG acc ttc ca ( SEQ ID No : 5 8 3 )
129	cct tct gga Cag gag cag ( SEQ ID No : 5 8 4 )
130	tac cag cag Aac gct tac g ( SEQ ID No : 5 8 5 )
131	g gag ggc cTg tgc gtg ( SEQ ID No : 5 8 6 )
132	g tac cag cGg gac gct t ( SEQ ID No : 5 8 7 )
133	c ggg tac cAg cag gac g ( SEQ ID No : 5 8 8 )
134	cag gac gcT tac gac gg ( SEQ ID No : 5 8 9 )
135	gtg cgg ttG gac agc ga ( SEQ ID No : 5 9 0 )
136	gag gac ggt Act cac acc ( SEQ ID No : 5 9 1 )
137	t ggc tgc Cac gtg ggg ( SEQ ID No : 5 9 2 )
138	ccg cgg gcA ccg tgg ( SEQ ID No : 5 9 3 )
139	cag aca cgg Cat gtg aag ( SEQ ID No : 5 9 4 )
140	g gcc cgt Tgg gcg gag ( SEQ ID No : 5 9 5 )
141	g gcc cgt Cgg gcg ga ( SEQ ID No : 5 9 6 )
142	tg gac gac Gcg cag ttc ( SEQ ID No : 5 9 7 )
143	cag ata atg Cat ggc tgc g ( SEQ ID No : 5 9 8 )
144	gag ggt ctC ccc aag cc ( SEQ ID No : 5 9 9 )
145	agg tat ttc Acc aca tcc g ( SEQ ID No : 6 0 0 )
146	at gtg aag gGc cac tca c ( SEQ ID No : 6 0 1 )
147	c acg gag ctT gtg gag ac ( SEQ ID No : 6 0 2 )
148	c ggg cgc Ctc ctc cg ( SEQ ID No : 6 0 3 )
149	g gat ggc aGg tgc gtg g ( SEQ ID No : 6 0 4 )
150	c ccc ccc aGg acg cat ( SEQ ID No : 6 0 5 )

Table 2-6

Probe No.	Base Sequence
151	ctg agc tcC tgg acc gc ( SEQ ID No : 6 0 6 )
152	g ata gag cGg gag ggg c ( SEQ ID No : 6 0 7 )
153	ccg tgg atG gag cag ga ( SEQ ID No : 6 0 8 )
154	c acg gac Gcc ccc aag ( SEQ ID No : 6 0 9 )
155	ag tgg gcg Tct gtg gtg ( SEQ ID No : 6 1 0 )
156	c ccc aag acG cat atg ac ( SEQ ID No : 6 1 1 )
157	g cag gag Agg ccg gag ( SEQ ID No : 6 1 2 )
158	gat tac atc Tcc ctg aac g ( SEQ ID No : 6 1 3 )
159	tc cgc aga Cac ctg gag ( SEQ ID No : 6 1 4 )
160	g aag tgg gTg gct gtg g ( SEQ ID No : 6 1 5 )
161	t ttc tac acT tcc gtg tcc ( SEQ ID No : 6 1 6 )
162	ac acc tcc Atg tcc cgg ( SEQ ID No : 6 1 7 )
163	c cgg cag Cac gcc tac ( SEQ ID No : 6 1 8 )
164	tat tgg gac Gag gag aca c ( SEQ ID No : 6 1 9 )
165	g gcg gcc cTt gtg gcg ( SEQ ID No : 6 2 0 )
166	c cgg cag gTc gcc tac ( SEQ ID No : 6 2 1 )
167	g gac ggg cAc ttc ctc c ( SEQ ID No : 6 2 2 )
168	g acc ctg cAc ggc tac t ( SEQ ID No : 6 2 3 )
169	cc atc cag aGg atg tat gg ( SEQ ID No : 6 2 4 )
170	c cag acc Agg gcg ggc ( SEQ ID No : 6 2 5 )
171	g cta ctc tTg ggg gcc c ( SEQ ID No : 6 2 6 )
172	g gac ctg gCg acc ctg ( SEQ ID No : 6 2 7 )
173	cac tca cag Gct gac cga ( SEQ ID No : 6 2 8 )
174	g gcg gcc Agt gtg gcg ( SEQ ID No : 6 2 9 )
175	gtg tcc cCg ccc ggc ( SEQ ID No : 6 3 0 )
176	t ctg ccc Gag ccc ctc ( SEQ ID No : 6 3 1 )

Table 3-1

Allele Number	Probe Number for Detection									
A*010101	0	1	2	3	4	5	6	7	8	
A*010102	9									
A*0102	10	11								
A*0103	12									
A*0106	13	14								
A*0107	15	16	17							
A*0108	18									
A*0109	19									
A*020101	20	21	22	23	24	25	26	27	28	
A*020102	29	30	31	21	22	23	24	32	33	34
A*020103	37									
A*020104	38									
A*020105	39									
A*020106	40									
A*020107	41	42								
A*020108	43									
A*020109	31	21	22	23	24	25	44	26	27	42
A*0202	45	42								
A*0203	20	46	47	48	27	28				
A*0204	20	21	22	24	25	26	27	28		
A*0205	45	28								
A*0206	20	49	21	22	23	24	25	26	27	28
A*0207	50									
A*0208	49	45								
A*0209	51									
A*0210	20	23	52	25	26	27	28			
A*0211	53	42	28							

Allele	Number	Probe Number for Detection																							
A*0212	20	25	54	27	28																				
A*0213	20	55	56	27	28																				
A*0214	45	26	28																						
A*0216	57	42	28																						
A*021701	20	58	24	25	26	27	28																		
A*021702	20	58	24	25	26	27	59																		
A*0218	60																								
A*0219	61	22	62	25	54	27																			
A*022001	29	63	30	21	22	23	24	32	33	34	35	25	26	27	36										
A*022002	64																								
A*0221	65																								
A*0222	20	21	22	23	24	25	44	27	28																
A*0224	29	30	31	21	22	23	24	32	33	35	25	26	27	36											
A*0225	46	66	26	27																					
A*0226	20	55	67	27	28																				
A*0227	22	68	69	27	36																				
A*0228	70	68	25	26	36																				
A*0229	71	68																							
A*0230	72																								
A*0231	73																								
A*0233	74																								
A*0234	31	75	76	22	23	24	25	44	26	27	42														
A*0235	31	75	22	23	24	32	33	34	35	25	26	27	36												
A*0236	29	30	31	21	22	23	24	32	33	34	35	25	26	27	36										
A*0237	22	68	25	54	27																				

[illegible]





Table 3-5

Allele Number	Probe Number for Detection									
A*240301	126	36	127							
A*240302	130									
A*2404	85	54	126	127						
A*2405	85	131	54	126	27					
A*2406	85	34	62	25	44	120	27			
A*2407	132	125	54	126	127					
A*2408	133	28								
A*2410	85	54	126	105	106					
A*2413	85	34	62	25	26	78	27			
A*2414	85	24	33	34	62	54	126	27		
A*2415	85	125	17	92	33	34	62	54	126	27
A*2417	85	125	17	58	104	34	62	54	126	27
A*2418	34	55	48	67	97					
A*2419	85	132	96	58	104	33	34	62	54	126
A*2420	85	125	17	58	104	33	34	62	54	126
A*2421	85	125	17	58	104	33	62	54	126	27
A*2422	44	36	127							
A*2423	85	54	126	27	134					
A*2424	91	58	34	80	81	122	78	27		
A*2425	123	54								
A*2426	135									
A*2427	136									
A*2428	85	61	17	58	104	33	34	62	54	126
A*2429	125	17	58	33	34	62	54	126	27	
A*2430	85	21	125	17	58	104	33	34	62	54
A*2431	137	25	54	27						
A*2432	138	34	54	27						
A*2433	62	25	54	27	42					
A*2434	53	54								

Table 3-6

Allele Number		Probe Number for Detection					
A*2435	139						
A*2437	140						
A*2438	141						
A*2501	138	142	28				
A*2502	91	138	142	28			
A*2503	138	143	47	48	106		
A*2504	138	47	56	106			
A*2601	90	48	142				
A*2602	144						
A*2603	21	61	48	142			
A*2604	145						
A*2605	16	48	142				
A*2606	146						
A*2607	31	48	142				
A*2608	56	142					
A*2609	147	131	143	47	27		
A*2610	34	131	143	47	48		
A*2612	131	143	66	44			
A*2613	91	147	131	143	47	48	
A*2614	49	90	147	148	55	48	
A*2615	149						
A*2616	10	90	147	131	143	47	48
A*2617	150						
A*2618	147	148	80	81	119		
A*29010101	151						
A*2902	152	36	28				
A*2903	152	28					
A*2904	153	80					
A*2905	152	56	36				

Table 3-7

Allele Number		Probe Number for Detection							
A*2906	122	154							
A*2907	152	58	122	36					
A*3001	10	15	155						
A*3002	11	15	156	27	36				
A*3003	11	156	27	36					
A*3004	11	25	36						
A*3006	157								
A*3007	86	156	27	36					
A*3008	49	15	155						
A*3009	11	81	122	36					
A*3010	158								
A*3011	10	155							
A*3012	15	156	27	36					
A*310102	15	121	159						
A*3102	84	53	104	147	121	80	122	36	
A*3103	53	160	80	122	36				
A*3104	160	159							
A*3105	15	53	104	147	121	80	122		
A*3106	15	53	104	121	80	122	36		
A*3107	15	125	147	121	81	122	36		
A*3108	161	85	125	147	121	122	36		
A*3109	162								
A*3201	125	122	163						
A*3202	54	163							
A*3203	125	164	80	122					
A*3204	138	97	165	166					
A*3205	167	125	122	163					
A*3206	138	25	26	27	36				
A*3207	10	138	80	81	122	27	36		

Table 3-8

Allele Number		Probe Number for Detection							
A*3301	168								
A*3303	90	121	159						
A*3304	169								
A*3305	170								
A*3306	171								
A*3401	172								
A*3402	47	67	27	36	173	174	175	28	
A*3403	160	55	67	27					
A*3404	70	47	67	36					
A*3405	176								
A*3601	177	79							
A*3602	178								
A*3603	179	177	79	36					
A*3604	105								
A*4301	114	142	28						
A*6601	91	96	48	142					
A*6602	57	175	28						
A*6603	47	57	180						
A*6604	47	181							
A*680101	49	91	104	44	182	28			
A*680102	183	91	104	44	182	28			
A*6802	184	28							
A*680301	183	104	44	182	28				
A*680302	183	35	44						
A*6804	90	53	68	36					
A*6805	183	21	35	44					
A*6806	91	89	68	25					
A*6807	91	185	68	25					



Table 4-1

Allele Number	Probe Number for Detection								
A*010101	0	1	2	3	4	5	6	7	
A*010102	8								
A*0102	9	10							
A*0103	11								
A*0106	12	13							
A*0107	14	15	16						
A*0108	17								
A*0109	18								
A*020101	19	20	21	22	23	24	25	13	26 27
A*020102	28	29	20	21	22	23	24	30 31 32 33	25 13 26 34
A*020103	35								
A*020104	36								
A*020105	37								
A*020106	38								
A*020107	39	27							
A*020108	40								
A*020109	20	21	22	23	24	25	12	13	26 27
A*0202	41	27							
A*0203	19	42	43	44	26	27			
A*0204	19	20	21	45	24	25	13	26	27
A*0205	46	41	27						
A*0206	19	46	20	21	22	23	24	25	13 26 27
A*0207	47								
A*0208	46	41							
A*0209	48								
A*0210	19	49	50	25	13	26	27		
A*0211	51	27							
A*0212	19	25	52	26	27				
A*0213	19	43	52	26	27				

Allele Number	Probe Number for Detection																							
A*0214	41	13	27																					
A*0216	53	27																						
A*021701	54																							
A*021702	19	55	24	25	13	26	56																	
A*0218	57																							
A*0219	58	22	59	25	52	26																		
A*022001	28	60	29	21	22	23	24	30	31	32	33	25	13	26	34									
A*022002	61																							
A*0221	62																							
A*0222	19	20	21	22	23	24	25	44	26	27														
A*0224	28	29	20	21	22	23	24	30	31	33	25	13	26	34										
A*0225	42	25	13	26																				
A*0226	19	43	13	26	27																			
A*0227	22	63	52	26	34																			
A*0228	64	63	25	13	34																			
A*0229	65	163																						
A*0230	66																							
A*0231	67																							
A*0233	68																							
A*0234	20	69	21	22	23	24	25	12	13	26	27													
A*0235	20	69	22	23	24	30	31	32	33	25	13	26	34											
A*0236	28	29	20	21	22	23	24	30	31	32	33	25	13	26										
A*0237	22	63	25	52	26																			
A*0238	63	42	52																					
A*0239	50	59	70	25	13	71	26	72	34															
A*0240	63	73	74	26	34																			
A*0241	46	28	29	20	21	22	23	24	75	32	33	25	13	26	34									
A*0242	76																							
A*0244	46	22	25	52	26	34																		



[illegible]

Table 4-4

Allele Number		Probe Number for Detection									
A*1103	96	97	52								
A*1104	46	69	58	52	26	72	34				
A*1105	98										
A*1106	69	21	58	52	72	34					
A*1107	99										
A*1108	46	69	58	43							
A*1109	100										
A*1110	46	82	58	52	34						
A*1111	101	58	52	34							
A*1112	46	69	58	16	52	72	34				
A*1113	102										
A*1114	95	103									
A*2301	104	13	71	105							
A*2302	77	32	73	74	44	106	26				
A*2303	31	107	73	13							
A*2304	77	32	73	74	13	71	26	72	34		
A*2305	108	13									
A*2306	109										
A*2309	13	71	105								
A*240201	77	110	52	111	105						
A*240202	77	110	16	55	93	31	32	52	111	26	
A*240203	112										
A*240204	113										
A*240301	111	34	105								
A*240302	77	52	111	26	72	34					
A*2404	77	52	111	105							
A*2405	77	114	52	111	26						
A*2406	77	32	59	25	44	106	26				
A*2407	69	110	52	111	105						

Table 4-5

Allele Number	Probe Number for Detection										
A*2408	115	116									
A*2410	77	52	111	72	34						
A*2413	77	32	59	25	13	71	26				
A*2414	77	24	31	32	59	52	111	26			
A*2415	77	110	16	83	31	32	59	52	111	26	
A*2417	77	110	16	55	93	32	59	52	111	26	
A*2418	32	43	12	13	88						
A*2419	77	69	58	55	93	31	32	59	52	111	26
A*2420	77	110	16	55	93	31	32	59	52	111	26
A*2421	77	110	16	55	93	31	59	52	111	26	
A*2422	117	34	105								
A*2423	77	52	111	26	118						
A*2424	69	55	32	73	74	13	71	26			
A*2425	108	52									
A*2426	119										
A*2427	120										
A*2428	77	58	16	55	93	31	32	59	52	111	26
A*2429	110	16	55	31	32	59	52	111	26		
A*2430	77	121	110	16	55	93	31	32	59	52	111
A*2431	122	25	52	26							
A*2432	123	32	52	26							
A*2433	59	25	52	26	27						
A*2434	124	52									
A*2435	125										
A*2437	126										
A*2438	127										
A*2501	123	128	129								
A*2502	69	123	128	129							
A*2503	123	42	43	44	34						

Table 4-6

Allele Number		Probe Number for Detection						
A*2504	123	43	52	34				
A*2601	82	44	128					
A*2602	130							
A*2603	21	58	44	128				
A*2604	131							
A*2605	15	44	128					
A*2606	132							
A*2607	20	44	128					
A*2608	52	128						
A*2609	133	114	42	43	26			
A*2610	32	114	42	43	44			
A*2612	114	42	25	44				
A*2613	69	133	114	42	43	44		
A*2614	46	82	133	134	43	44		
A*2615	135							
A*2616	9	82	133	114	42	43	44	
A*2617	136							
A*2618	133	134	73	74	44			
A*29010101	137							
A*2902	138	34	129					
A*2903	138	129						
A*2904	139	73						
A*2905	138	52	34					
A*2906	138	13	34					
A*2907	138	55	13	34				
A*3001	9	14	140					
A*3002	10	14	141	26	34			
A*3003	10	141	26	34				
A*3004	10	25	34					

Table 4-7

Allele Number		Probe Number for Detection							
A*3006	142								
A*3007	78	141	26	34					
A*3008	46	14	140						
A*3009	10	74	13	34					
A*3010	143								
A*3011	9	140							
A*3012	14	141	26	34					
A*310102	14	107	144						
A*3102	20	51	93	133	107	73	13	34	
A*3103	51	134	73	13	34				
A*3104	134	144							
A*3105	14	51	93	133	107	73	13		
A*3106	14	51	93	107	73	13	34		
A*3107	14	110	133	107	74	13	34		
A*3108	145	77	110	133	107	13	34		
A*3109	146								
A*3201	123	13	147	129					
A*3202	123	52	147	129					
A*3203	110	148	73	13					
A*3204	123	88	149	34					
A*3205	78	123	13	147	129				
A*3206	123	25	13	26	34				
A*3207	9	123	73	74	13	26	34		
A*3301	150								
A*3303	82	107	144						
A*3304	151								
A*3305	152								
A*3306	153								
A*3401	133	43	44	26	34	154	155	129	

Table 4-8

Allele Number		Probe Number for Detection						
A#3402	43	13	26	34	154	156	155	129
A#3403	134	43	13	26				
A#3404	157	43	13	34				
A#3405	158							
A#3601	26	72						
A#3602	88							
A#3603	83	26	72	34				
A#3604	72							
A#4301	101	128	129					
A#6601	69	58	44	128				
A#6602	53	155	129					
A#6603	43	53	34					
A#6604	43	159						
A#680101	46	69	93	44	160	27		
A#680102	161	69	93	44	160	27		
A#6802	162	27						
A#680301	161	93	44	160	27			
A#680302	161	33	44					
A#6804	82	51	63	34				
A#6805	161	21	33	44				
A#6806	69	81	63	25				
A#6807	69	163	63	25				
A#6808	89	160	27					
A#6809	161	52						
A#6810	46	164	69	93	25	44	26	34
A#6812	161	69	44					
A#6813	46	69	93	44	160			
A#6814	46	164	69	93	25	44	26	34
A#6815	162	82	21					

[illegible]

(Example 3)

Probes for identification of HLA-B allele

Extraction of DNA from 1 ml of human blood was performed using GFX Genomic Blood DNA Purification Kit from Amersham Biosciences in the same manner as in Example 1.

Next, quantitative PCR was carried out in the same manner as in Example 1 except that probes in the probe list B1 were used and 3  $\mu$ l of the mixed primers consisting of 1  $\mu$ l each of respective solutions of the following primers (10 pmol/ $\mu$ l):

CTGAGCTCTTCCTCCTACACA (SEQ ID NO:518)

TCCTTCCCGTTCTCCAGGT (SEQ ID NO:519)

AGGTCTCGGTCAGGGCCA (SEQ ID NO:520)

After PCR amplification, the sample was identified being B\*520101, referring to Amp Plot and Dissociation curves on a display of 5700 software and the allele-probe list B1 (described later).

(Example 4)

Extraction of DNA from 1 ml of human blood was performed in the same way as in Example 1. PCR of human HLA-B was then performed in the same manner as in Example 2 except that 2  $\mu$ l of the mixed primer consisting of 1  $\mu$ l each of the respective solutions of the following primers at 10 pmol/ $\mu$ l and 13  $\mu$ l of ultra pure water used:

CTGAGCTCTTCCTCCTACACA (SEQ ID NO:518)



GCTCCCACTCCATGAGGTATTTTC (SEQ ID NO:521).

At the same time, a DNA microarray was prepared to identify the allele in the specimen described above in the same manner as in Example 2, except that probes in the probe list B2 were to form the probe dots respectively.

Then, hybridization was performed using the above specimen and the prepared DNA microarray in the same manner as in Example 2. Fluorometry measurement was conducted with GenePix4000B (Axon). Referring to the allele-probe list B2 (described later), the sample was identified as B\*520101.

#### Allele list

15 B\*070201

atgctgggtcatggcgccccgaaccgtccctctgctgctctcgggcgccccggccctgaccgagacctgggcccggctccca  
ctccatgaggatattctacacctccGtgtcccgccccggcggggagccccgcttcatctcagtgggctacgtggacg  
acaccagttcgtgaggttcgacagcgacgccgagtcgagagaggagccgccccggcgccgtggatagagcaggagggg  
ccggaglatgggaccggaacacacagatctacaaggcccaggcacagactgaccgagagagccgtcggaacctgcgcgg  
20 ctactacaaccagagcgaggccgggtctcacacctccagagCatgtacggctgcgacgtggggccggacgggcgcctcc  
tccgcgggCatgaccagTaccctacgacggcaaggattacatcgccctgaacgaggacctgcgtctctggaccgccg  
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cccaccaccccatctctgaccatgaggccacctgagggtgctgggccccgggtttctacctgaggagatcacactgacc  
25 tggcagcgggatggcgaggaccaaactcaggacactgagcttggagaccagaccagcaggagatagaacctccagaa  
gtgggcagctgtgggtgggtgcttctggagaagagcagagatacacatgccatgtacagcatgaggggctgccgaagcccc  
tcacctgagatgggagccgtcttccagctccaccgtccccatcgctgggcattgttgcctggcctggctgtcttANNgca

gttgggtcatcggagcgtggcgcgtgctgtgatgtgtaggaggaagagttcaggtagga (SEQ ID NO:522)

B\*070202

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ctccatgaggtaattctacacctccgtgtcccgcccgcccgccggggagccccgttcatctcagtagggctacgtggacg  
5 acaccagttcgtgaggttcgacagcgacgcccgagtcgagagaggagcccgccggcgccgtggatagagcaggagggg  
ccggagtaattgggaccggaacacacagatctacaaggcccaggcacagactgaccgagagagccitcggaacctgcgcgg  
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gacacggcggctcagatcaccagcgcaagtgggaggcgcccgtagggcggagcagcggagagccctaccitggagggcga  
10 gtgcgtggagtggctccgcagGtacctggagaacgggaaggacaagctggagcgcgtgacccccaaagacacacgtga  
cccaccacccatctctgaccatgaggccacctgaggtgtcgggcccgtgggtttctacctgaggagatcacactgacc  
tggcagcgggatggcgaggaccaaactcaggacactgagctgtggagaccagaccagcaggagatagaaccttccagaa  
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taccttgagatgggagccgtcttccagttccacctccccatcgtgggcatgtgtgtcgtggcctggctgtccta... gca  
15 gtgtgggtcatcggagcgtggcgcgtgctgtgatgtgtaggaggaagagtt (SEQ ID NO:523)

B\*070203

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gtggacgacacGcagttcgtgaggttcgacagcgacgcccgagtcgagagaggagcccgccggcgccgtggatagagca  
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20 tgcgcggctactacaaccagagcgaggccgggtctcacacctccagagcaigtacggctgcgacgtggggccggacggg  
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NO:524)

25 B\*0703

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10 taccctgagatgggagccgtcttccagtcaccgtccccatcgltgggcatgtgtgctggcctggctgtccta... gca  
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B\*0704

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15 acaccagatcgtaggttcgacagcgacgccgcgagtcgagagaggagccgcgggcgccgtggatagagcaggagggg  
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gacacggcggctcagatcaccagcgcaagtgggaggcgcccgtagggcggagcaggaCagagccctaccitggaggcgga  
20 gtgcgtggagtggctccgcagatacctggagaacgggaaggacaagctggagcgcgctgacccccaaagacacacgtga  
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taccctgagatgggagccgtcttccagtcaccgtccccatcgltgggcatgtgtgctggcctggctgtccta... gca  
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B\*0705

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B\*0706  
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25 tcacctgagatgggagccgtcttccagtcaccgtccccatcgtgggcatgtgtgctggccctggctgtccta... gca  
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B\*0707

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B\*0708
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B\*0709
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B\*0710

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g t g g a c g a c a c c c a g t t c g t g a g g t t c g a c a g c g a c g c c g c g a g t c c g a g a g a g g a g c c g c g g g c g c c g t g g a t a g a g c a  
g g a g g g c c g g a g t a t t g g g a c c g g a a c a c a c a g a t c t c a a g g c c c a g g c a c a g a c t g a c c g a g a g a c c t g c g g a a c c  
t g c g c g g c t a c t a c a a c c a g a g c g a g g c c g g g t c t c a c a c c t c c a g a g c a t g t a c g g c t g c g a c g t g g g c c g g a c g g g  
5 c g c c t c c t c c g c g g g c a t g a c c a g t a c g c c t a c g a c g g c a a g g a t t a c a t c g c c c t g a a c a g g a c c t g c g c t c c t g g a c  
c g c c g c g g a c a c g g c g g c t c a g a t c a c c c a g c g c a a g t g g g a g g c g g c c c g t g a g g c g g a g c a g c g g a g a g c c t a c c t g g  
a g g g c g a g t g c g t g g a g t g g c t c c g c a g a t a c c t g g a g a a c g g g a a g g a c a a g c t g g a g c g c g c t g (SEQ ID  
NO:532)

B\*0711

10 g c t c c a c t c c a t g a g g t a t t t c t a c a c c t c c g t g t c c c g g c c c g g c c g g g g a g c c c c g c t t c a t c t c a g t g g g c t a c  
g t g g a c g a c a c c c a g t t c g t g a g g t t c g a c a g c g a c g c c g c g a g t c c g a g a g a g g a g c c g c g g g c g c c g t g g a t a g a g c a  
g g a g g g c c g g a g t a t t g g g a c c g g a a c a c a c a g a t c t a c a a g g c c c a g g c a c a g a c t g a c c g a g a g a c c t g c g g a a c c  
t g c g c g g c t a c t a c a a c c a g a g c g a g g c c g g g t c t c a c a c c t c c a g a g c a t g t a c g g c t g c g a c g t g g g c c g g a c g g g  
c g c c t c c t c c g c g g g c a t g a c c a g t C g c c t a c g a c g g c a a g g a t t a c a t c g c c c t g a a c a g g a c c t g c g c t c c t g g a c  
15 c g c c g c g g a c a c g g c g g c t c a g a t c a c c c a g c g c a a g t g g g a g g c g g c c c g t g a g g c g g a g c a g c g g a g a g c c t a c c t g g  
a g g g c g a g t g c g t g g a g t g g c t c c g c a g a t a c c t g g a g a a c g g g a a g g a c a a g c t g g a g c g c g c t g (SEQ ID  
NO:533)

B\*0712

20 g c t c c a c t c c a t g a g g t a t t t c t a c a c c t c c g t g t c c c g g c c c g g c c g g g g a g c c c c g c t t c a t c t c a g t g g g c t a c  
g t g g a c g a c a c c c a g t t c g t g a g g t t c g a c a g c g a c g c c g c g a g t c c g a g a g a g g a g c c g c g g g c g c c g t g g a t a g a g c a  
g g a g g g c c g g a g t a t t g g g a c c g g a a c a c a c a g a t c t a c a a g g c c c a g g c a c a g a c t g a c c g a g a g a c c t g c g g a a c c  
t g c g c g g c t a c t a c a a c c a g a g c g a g g c c g g g t c t c a c a t c A t c c a g a g g a t g t a T g g c t g c g a c C t g g g g c c C g a c g g g  
c g c c t c c t c c g c g g g c a t g a c c a g t a c g c c t a c g a c g g c a a g g a t t a c a t c g c c c t g a a c a g g a c c t g c g c t c c t g g a c  
c g c c g c g g a c a c g g c g g c t c a g a t c a c c c a g c g c a a g t g g g a g g c g g c c c g t g a g g c g g a g c a g c g g a g a g c c t a c c t g g  
25 a g g g c g a g t g c g t g g a g t g g c t c c g c a g a t a c c t g g a g a a c g g g a a g g a c a a g c t g g a g c g c g c t g (SEQ ID  
NO:534)

B\*0713

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B\*0714

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5 B\*0719

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B\*0720

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5 B\*0722

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B\*0723

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5 B\*0728

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5 B\*0731

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B\*0810

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B\*1401

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20 B\*1527

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25 B\*1533

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A g a g g g g c c g g a g t a t t g g g a c c g g a a c a c a c a g a t c t C c a a g a c c a a c a c a c a g a c t T a c c g a g a g a g c c t g c g g a a c c  
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a g g g c a c g t g c g t g g a g t g g c t c c g c a g a C a c c t g g a g a a c g g g a a g g a g a c g c t g c a g c g c g c g (SEQ ID  
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c g c c t c c t c c g c g g g c a t g a c c a g t c c g c c t a c g a c g g c a a g g a t t a c a t c g c c c t g a a G a g a g g a c c t g a g c t c c t g g a c  
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B\*3506

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B\*3513



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B\*3514
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5 B\*3516

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 g g a g g g c c g g a g t a t t g g g a c c g g g a G a c a c a g a t c t T c a a g a c c a a c A c a c a g a c t T a c c g a g a g a g c c t g c g g a a c c  
 t g c g c g g c t a c t a c a a c c a g a g c g a g g c c g g g t c t c a c a t c A t c c a g a g c a t g t a c g g c t g c g a c t g g g g c c G a c g g g  
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B\*3517

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 NO:725)

B\*3518

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5 B\*3519

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B\*3520

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NO: 728)

B\*3521

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5 B\*3522

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B\*3524

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B\*3527

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5 B\*3528

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B\*3529

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5 B\*3531

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10 B\*3534

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NO:783)



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10 B\*3537

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NO:785)g

B\*3538

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B\*3541

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B\*3542

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[illegible]

B\*3543

[illegible]

B\*3544

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NO:791)

5 B\*3545

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B\*3701

[illegible]

B\*3702

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15 B\*3704

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B\*3801

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25 B\*380201

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20 NO:799)  
B\*3803  
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B\*3804

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B\*3806

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B\*4007

10 atgcggtcacggcaccgccgaaccgtctctcgtcgtctcggcgccctggccctgaccgagacctgggcccgtccca  
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15 tccgcgggcatAaccaglacgccacgacggcaaggattacatcgccctgaacgaggacctgcgcctctggaccgcccgcg  
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B\*4008

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- ciactacaaccagagcgaggccgggtctcacacccctccagagcaigtacggctgcgacgtggggccggacgggcgccctcc  
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- 10 B\*4009  
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NO: 847)
- B\*4010  
20 gtctccactccatgaggatatttctacaccgccatgtcccgcccgccggcgaggagccccgccttcatcCagtgggcttac  
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NO: 848)

B\*4011

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NO:849)

10 B\*4012

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B\*4013

20 atgcggtcacggcgccccgaaccctctctctgtgtcttcgggggagtgccctgaccgagacctgggtTggctccca  
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B\*401401

gc tccca tcca tga ggtat ttcCac accgcca tgc tccggcccggccg cggggagccccgc t tca taccgtgggc tac  
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5 tgcgcggctactacaaccagagcgaggccgggtctcacaccc tccagagga tgtacggctgcgacgtggggccggacggg  
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cgcgccggacacCgcggctcagatcaccagcgcaagtgggaggcgccccgtTggcggagcagcTgagagcc tacc tgg  
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NO: 852)

10 B\*401402

gc tccca tcca tga ggtat ttcCac accgcca tgc tccggcccggccg cggggagccccgc t tca taccgtgggc tac  
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NO: 853)

B\*4015

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tgcgcggctactacaaccagagcgaggccgggtctcacaccc tccagagca tgtacggctgcgacgtggggccggacggg  
20 cgcc tcc tccg cgggca tAaccagtacgcttacgacggcaaggattacatcgccctgaacgaggacctgcgctcctggac  
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NO: 854)

B\*4016

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 5 ccggaglatlgggaccgggaGacacagatctCcaagaccaacacacagactTaccgagagagcctgcggaaccigcgcgg  
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10 B\*4018

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 NO:856)

B\*4019

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 25 cgcccgggacacggcggtcagatcaccagcgcaagtgggaggcgcccgctgTggcgagcagcTgagagcctacctgg  
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 NO:857)

B\*4020

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 5 ccggagtatgggaccgggaGacacagatctCcaagaccaacacacagacTaccgagagagccctgcggaacctgcgcgg  
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10 B\*4021

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 NO:859)

B\*4023

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B\*4024

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NO:861)

10 B\*4025

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NO:862)

B\*4026

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NO:863)



B\*4027

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NO:864)

10 B\*4028

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NO:865)

B\*4029

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NO:866)

B\*4030

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5 tgcgcggct|act|acaaccagagcgaggccgggt|ct|cacat|catccagG|tga|gt|atggc|tgcgacgt|ggggccggacggg  
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cgccg|cgacacggcggt|cagat|c|cccagcgcaagi|Tggaggcgcccg|gt|ggcggagcagc|Tgagagcc|tacc|tgg  
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NO: 867)

10 B\*4031

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NO: 868)

B\*4032

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NO: 869)

B\*4033

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NO: 870)

10 B\*4034

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B\*4035

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B\*4036

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10 B\*4037

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NO: 874)

B\*4038

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NO: 875)



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NO:888)

B\*4204

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10 NO:889)

B\*440201

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B\*440202

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B\*440302

[illegible]

B\*4404

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15, B\*4406

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B\*4407

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g t t g t g g t c a t c g g a g c t g t g g t c g c t g c t g t a t g t g t a g g a g a a g a g C t a g g t g g a ( S E Q I D N O : 9 0 0 )  
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20 N O : 9 0 1 )  
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B\*4424

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B\*4425

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NO:914)

B\*4426

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B\*4428

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B\*5109

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NO: 962)

B\*5110

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5 B\*5112

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 NO: 964)

B\*511301

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 NO: 965)

B\*511302

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NO:966)

5 B\*5114

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NO:967)

B\*5115

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NO:968)

B\*5116

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NO:969)

5 B\*5117

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NO:970)

B\*5118

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NO:971)

B\*5119

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NO:972)

5 B\*5120

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NO:973)

B\*5121

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NO:974)

B\*5122

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NO:975)

5 B\*5123

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NO:976)

B\*5124

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NO:977)

B\*5126

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NO:978)

5 B\*5128

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NO:979)

B\*5129

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25 aagccccctacccctgagatggg (SEQ ID NO:980)

B\*5130

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 B\*5131  
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 NO:982)  
 20 B\*5132  
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5 B\*5133

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B\*5134

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NO: 1052)

## B\*7804

25 gctccca tccatgaggtat tctacaccgccatgtcccgccccggccgaggagccccgc t tcatgcag tgggctac  
gtggacgacacccag t tctgaggt tgcacagcgacgcccgag tccgaggacggagccccgggcgccatggatagagca  
ggagggggccggagtat tgggaccggaacacacagat tctCaagaccaacacacagact taccgagagagcc tgcggaacc  
tgcgcggctactacaaccagagcgaggccgggtctcacact tggcagacgatgtatTggctgcgacgtggggccggacggg

cgccctccctccgcgggcataaaccagtacgcctacgacggcaaAgattacaicgccctgaacgaggacctgAgctccctggac  
cgcgcgggacaccgcggctcagatcacccagcgcaagtgggaggcgggcccgTggcgggagcagTgagagccctacctgg  
agggccTgtgcgtggagtggctccgcagatacctggagaacgggaaggagacgctgcagcgcgcggaacccccaaagaca  
cacgtgaccaccaccccGtctctgacctgaggccacctgaggctgctgggccctgggcttctaccttgcggagatcac  
5 actgacctggcagcgggatggcgaggaccaaactcaggacactgagcttctggagaccagaccagcaggagatagaacct  
tccagaagtgggcagctgtgggtggcttctggagaagagcagagatcacatgccatgtacagcatgaggggctgccg  
aagccccctacccctgagatggg (SEQ ID NO:1056)

B\*7805

gtccccactccatgaggatcttctacaccgccaatgccccggccggcgcggggagccccgcttcatcgcatgggcctac  
10 gtggacgacaccagttcgtgaggttcgacagcgacgccgcgagctccgaggaCggagccCggggcgccatggatagagca  
ggaggggcccggaglatgggaccgggaGacacagatctCcaagaccaacacacagactTaccgagagagccctgcggaacc  
tgcgcggctactacaaccagagcgaggccgggtctcacacttggcagaCgatgtTggctgcgacgtggggccggacggg  
cgccctccctccgcgggcataaaccagtacgcctacgacggcaaAgattacatcgccctgaacgaggacctgagctccctggac  
cgcgcgggacaccgcggctcagatcacccagcgcaagtgggaggcgggcccgTgagcgggagcagTgagagccctacctgg  
15 agggccTgtgcgtggagtggctccgcagaCacctggagaacgggaaggagacgctgcagcgcgcg (SEQ ID  
NO:1054)

B\*8101

atgttggtcatggcgccccgaaccgtccctccctgctgctctggggggcagtgggccctgaccgagacctgggcccgtccca  
ctccatgaggatcttctacacctccgtgtcccgcccgcccgcggggagccccgcttcatctcagtgggctacgtggacg  
20 acaccagttcgtgaggttcgacagcgacgccgcgagtcgagagaggagccgcggcgccgtggatagagcaggagggg  
ccggaglatgggaccgggaacacacagatctacaaggcccaggcacagactgaccgagagagccctgcggaacctgcgcgg  
ctactacaaccagagcgaggccgggtctcacacctccagagcatgtacggctgcgacgtggggccggacgggcgccctcc  
tccgcgggcataaaccagtacgcctacgacggcaaggattacatcgcccgaacgaggacctgcgctccctggaccgccgcg  
gacacggcggtcagatctccagcgcaagtgggaggcgggcccgTgtggcgggagcagctgagagccctacctggaggcgga  
25 gtgcgtggagtggctccgcagatacctggagaacgggaaggacaagctggagcgcgctgacccccaaagacacacgtga  
cccaccaccccatctctgacctgaggccacctgaggctgctgggccctgggttcttaccttgcggagatcacactgacc  
tggcagcgggatggcgaggaccaaactcaggacactgagcttctggagaccagaccagcaggagatagaaccttccagaa

gtggacagctgiggtggccttctggagaagagcagagatcacatgccatgtacagcatgaggggctgccgaagcccc  
tcacccigagatgggagccgtcttcccagtcacccgtcccatcgtgggcattgttgcctggcctgtctta... gca  
gttgtggcatcggagctgiggtcgctgctgtgatgtgtaggaggaagagctcTggtgga (SEQ ID NO:1055)

B\*8201

5 gctcccactccatgaggtatttctacaccgctatgtcccgcccgccgaggagccccgcttcaatcagtgggctac  
giggacgacacgcagttcgtaggttcgacagcgacgccgcgagtcgagagaggagccgaggcgccgtggatagagca  
ggagggcgccgagttatgggaccggaacacacagatctacaaggcccaggcacagactgaccgagagagccctgcggaacc  
tgcgcggctactacaaccagagcgaggccgggtctcacacccctcagaggatgtttggctgcgacctggggcccgacggg  
cgcttctccgcgggcataaccagttagcctacgacggcaaggattacatcgcccgaacgaggacctgagctcttgga  
10 cgcggcgacacccggtcagatcaccagcgcaagtgaggagcgcccgctgtggcgagcaggacagagcctacctgg  
aggAccgtgtcgtggagtcgctccgcagataccitggagaacgggaaggagacgctgcagcgcgcgg

(SEQ ID NO:1056)

B\*8202

atgcggtcacggcaccccgaaacctctctctgctcttggggggccctggccctgaccgagacctgggctggctccca  
15 ctccatgaggtatttctacaccgctatgtcccgcccgccgaggagccccgcttcaatcagtgggctacgtggacg  
acacgcagttcgttaggttcgacagcgacgccgcgagtcgagagaggagccgaggcgccgtggatagagcaggagggg  
ccgagttatgggaccggaacacacagatctacaaggcccaggcacagactgaccgagagagccctgcggaacctgcgcgg  
ctactacaaccagagcgaggccgggtctcacacccctcagaggatgtttggctgcgacctggggcccgacggcgccctcc  
tccgcgggcataaccagttagcctacgacggcaaggattacatcgcccgaacgaggacctgagctcttgaccgcggcg  
20 gacaccgcggctcagatcaccagcgcaagtgaggagcgcccgctgtggcgagcaggacagagcctacctggaggccct  
gtgcgtggagtcgctccgcagataccitggagaacgggaaggagacgctgcagcgcgcggacccccaaagacacatgtga  
cccaccacccatctctgacctgaggccacctgaggctcggggccctgggcttctacctgaggagatcacactgacc  
tggcagcgggatggcgaggacaaactcaggacaccgagctgtggagaccagaccagcaggagatagaacctccagaa  
gtgggcagctgtgtggccttctggagaagagcagagatcacatgccatgtacagcatgaggggctgccgaagcccc  
25 tcacccigagatgggagccatcttcccagtcacccatcccatcgtgggcattgttgcctggcctgtctta... gca  
gttgtggcatcggagctgtgtTgctactgtgatgtgtaggaggaagagctcaggtgga (SEQ ID NO:1057)

B\*8301

atgcgggtcacggcgccccgaacccctccctcctgctgctctggggggcaGiggccctgaccgagacctgggcccgtccca  
ctccatgagglattttctacaccgccAtgtcccgcccgcccggggagccccgcttcatctcagtgggctacgtggacg  
acacccagttcgtgaggttcgacagcgacgccgcgagtcgagagaggagcgccggcgccgtggatagagcaggagggg  
ccggagtattgggaccggaacacacagatctacaaggcccaggcacagactgaccgagagagcctgcggaacctgcgcgg  
5 ctactacaaccagagcgaggccgggtctcacatcAtccagaggatgtacggctgcgacgtggggccggacggcgccctcc  
tccgctggTatgaccagGacgcttacgacggcaaggattacatcgccctgaacgaggacctgagctcctggaccgcggcg  
gacaccgcggctcagatcaccagcgcaagtgaggaggcgcccgctgtggcgagcaggaCagagccctacgtggaggccT  
gtgcgtggagtCgctccgcagataacctggagaacgggaaggagacgctgcagcgcgCg (SEQ ID NO:1058)

10           The following Tables 5-1 to 5-9 show Probe list  
B1, and Tables 6-1 to 6-8 show Probe list B2. The  
Allele-probe list is shown in Tables 7 and 8.

Table 5-1

Probe No.	Base Sequence
0	agg tat ttc tac acc tcc G ( SEQ ID No: 1)
1	ct cac acc ctc cag agC ( SEQ ID No: 2)
2	gc ctc ctc cgc ggg C ( SEQ ID No: 3)
3	c cgc ggg cat gac cag T ( SEQ ID No: 4)
4	gt gag gcg gag cag cG ( SEQ ID No: 5)
5	t gag gcg gag cag cgG ( SEQ ID No: 6)
6	gcc tac ctg gag ggc gA ( SEQ ID No: 7)
7	ggc gag tgc gtg gag tG ( SEQ ID No: 8)
8	c ggg aag gac aag ctg G ( SEQ ID No: 9)
9	g gag tgg ctc cgc agG ( SEQ ID No: 10)
10	gc tac gtg gac gac acG ( SEQ ID No: 11)
11	a cag atc tac aag acc aac A ( SEQ ID No: 12)
12	gt gag gcg gag cag gaC ( SEQ ID No: 13)
13	c ctc ctc cgc ggg cat A ( SEQ ID No: 14)
14	cg tct tcc cag tcc acc A ( SEQ ID No: 15)
15	ct cac acc ctc cag agG ( SEQ ID No: 16)
16	ac cgg aac aca cag atc tT ( SEQ ID No: 17)
17	a cag atc ttc aag acc aac A ( SEQ ID No: 18)
18	cgc ggg cat gac cag tC ( SEQ ID No: 19)
19	c cgg aac aca cag atc tG ( SEQ ID No: 20)
20	ca cag act gac cga gag aA ( SEQ ID No: 21)
21	g gcc ggg tct cac atc A ( SEQ ID No: 22)
22	ac atc atc cag agg atg taT ( SEQ ID No: 23)
23	gg atg tat ggc tgc gac C ( SEQ ID No: 24)
24	c tgc gac ctg ggg ccC ( SEQ ID No: 25)
25	ag aca cag aag tac aag cG ( SEQ ID No: 26)
26	c aag cgc cag gca cag G ( SEQ ID No: 27)
27	gca cag gct gac cga gT ( SEQ ID No: 28)
28	gag gcc ggg tct cac aT ( SEQ ID No: 29)
29	g tct cac atc atc cag agG ( SEQ ID No: 30)
30	cgc ctc ctc cgc ggg T ( SEQ ID No: 31)

Table 5-2

Probe No.	Base Sequence
31	c aag gcc cag gca cag G (SEQ ID No:32)
32	c aag acc aac aca cag act T (SEQ ID No:33)
33	cgc ggg tat gac cag tC (SEQ ID No:34)
34	gcc tac ctg gag ggc aC (SEQ ID No:35)
35	ctg gag aac ggg aag gaG (SEQ ID No:36)
36	g acg ctg gag cgc gcG (SEQ ID No:37)
37	gcc tac ctg gag ggc cT (SEQ ID No:38)
38	ggc ctg tgc gtg gag tC (SEQ ID No:39)
39	c ggc cgc ggg gag cT (SEQ ID No:40)
40	tcc tgg acc gcc gcg A (SEQ ID No:41)
41	cgg aac ctg cgc ggc C (SEQ ID No:42)
42	gcc tac ctg gag ggc C (SEQ ID No:43)
43	gg gag gcg gcc cgt gT (SEQ ID No:44)
44	gt gtg gcg gag cag gaC (SEQ ID No:45)
45	cgt gag gcg gag cag cT (SEQ ID No:46)
46	c cgg aac aca cag atc tC (SEQ ID No:47)
47	ca cag act tac cga gag G (SEQ ID No:48)
48	ctg cgg acc ctg ctc C (SEQ ID No:49)
49	c cgc ggg tat gac cag G (SEQ ID No:50)
50	cac tcc atg agg tat ttc G (SEQ ID No:51)
51	gg tat ttc gac acc gcc A (SEQ ID No:52)
52	cg aga gag gag ccg cC (SEQ ID No:53)
53	a gcc tac ctg gag ggc A (SEQ ID No:54)
54	g atg tgt agg agg aag agC (SEQ ID No:55)
55	ctg cgc acc gcg ctc C (SEQ ID No:56)
56	c cga gag aac ctg cgg aT (SEQ ID No:57)
57	gag aac ctg cgg atc gC (SEQ ID No:58)
58	ctg cgg atc gcg ctc C (SEQ ID No:59)
59	c acg ctg gag cgc gcG (SEQ ID No:60)
60	g gac cgg aac aca cag aC (SEQ ID No:61)

Table 5-3

Probe No.	Base Sequence
61	c act tgg cag acg atg taT ( SEQ ID No:62)
62	g gag tat tgg gac cgg G ( SEQ ID No:63)
63	c cgg gac aca cag atc tT ( SEQ ID No:64)
64	cgt gtg gcg gag cag cT ( SEQ ID No:65)
65	cgc ggg tac cac cag G ( SEQ ID No:66)
66	c aca cag act gac cga gT ( SEQ ID No:67)
67	ttc aag acc aac aca cag G ( SEQ ID No:68)
68	c cgg gag aca cag atc tC ( SEQ ID No:69)
69	g tgc tgg gcc ctg ggC ( SEQ ID No:70)
70	g gct cag atc acc cag cT ( SEQ ID No:71)
71	g tct cac act tgg cag aC ( SEQ ID No:72)
72	cgc ggg cat aac cag ttA ( SEQ ID No:73)
73	cg atg tat ggc tgc gac C ( SEQ ID No:74)
74	tgg gag cca tct tcc caA ( SEQ ID No:75)
75	gag cag ctg aga gcc tG ( SEQ ID No:76)
76	gg tct cac acc ctc cag T ( SEQ ID No:77)
77	cc aga cca gca gga gaC ( SEQ ID No:78)
78	cc ctg aga tgg gag caA ( SEQ ID No:79)
79	c atg agg tat ttc tac acc G ( SEQ ID No:80)
80	c tcc cac tcc atg agg C ( SEQ ID No:81)
81	g cag gag ggg ccg gaA ( SEQ ID No:82)
82	g gag tgg ctc cgc aga C ( SEQ ID No:83)
83	g acg ctg cag cgc gcG ( SEQ ID No:84)
84	c acc ctc cag agg atg taT ( SEQ ID No:85)
85	tc ctg ctg ctc tcg ggA ( SEQ ID No:86)
86	gcg ccc cgg gcg ccA ( SEQ ID No:87)
87	gag tat tgg gac cgg gaG ( SEQ ID No:88)
88	c cgt gag gcg gag cag T ( SEQ ID No:89)
89	gac caa act cag gac acC ( SEQ ID No:90)
90	cc gcc tac gac ggc aaA ( SEQ ID No:91)



Table 5-4

Probe No.	Base Sequence
91	g agc tcc tgg acc gcG (SEQ ID No:92)
92	g gat tac atc gcc ctg aaT (SEQ ID No:93)
93	c gac acg cag ttc gtg C (SEQ ID No:94)
94	cag atc tcc aag acc aac A (SEQ ID No:95)
95	c gga gct gtg gtc gct A (SEQ ID No:96)
96	c acc ctc cag agg atg tT (SEQ ID No:97)
97	tac gcc tac gac ggc aaA (SEQ ID No:98)
98	cag atc tgc aag acc aac A (SEQ ID No:99)
99	cg agt ccg agg atg gcT (SEQ ID No:100)
100	g ggc ctg tgc gtg gaC (SEQ ID No:101)
101	gg gcc ggc tcc cac tT (SEQ ID No:102)
102	ac atg aag gcc tcc gcG (SEQ ID No:103)
103	gca gct gtg gtg gtg cT (SEQ ID No:104)
104	gtg acc cac cac ccc G (SEQ ID No:105)
105	g tat tgg gac cgg gag aT (SEQ ID No:106)
106	gcg agt ccg agg atg gC (SEQ ID No:107)
107	c acc ctc cag agg atg tC (SEQ ID No:108)
108	gg acc gcc gcg gac aA (SEQ ID No:109)
109	g atg tac ggc tgc gac C (SEQ ID No:110)
110	g tct cac acc ctc cag aC (SEQ ID No:111)
111	ct cac acc ctc cag acG (SEQ ID No:112)
112	ac cga gag aac ctg cgC (SEQ ID No:113)
113	c ggg aag gag acg ctg C (SEQ ID No:114)
114	cc ctg aac gag gac ctg A (SEQ ID No:115)
115	g gag ccc cgc ttc atc G (SEQ ID No:116)
116	agg tat ttc tac acc gcc A (SEQ ID No:117)
117	t ccg agg atg gcg ccC (SEQ ID No:118)
118	g ttc gac agc gac gcc A (SEQ ID No:119)
119	gag ccg cgg gcg ccA (SEQ ID No:120)
120	g gcg gag cag ctg aga A (SEQ ID No:121)

Table 5-5

Probe No.	Base Sequence
121	a acc tac ctg gag ggc C (SEQ ID No:122)
122	acc tac ctg gag ggc cT (SEQ ID No:123)
123	c tcc aag acc aac aca cG (SEQ ID No:124)
124	c tac gtg gac gac acg cT (SEQ ID No:125)
125	c cgg gag aca cag atc tT (SEQ ID No:126)
126	ac aca cag act tac cga gT (SEQ ID No:127)
127	ca cag act tac cga gtg aA (SEQ ID No:128)
128	c cgc ggg cat aac cag tT (SEQ ID No:129)
129	cc cag ttc gtg agg ttc A (SEQ ID No:130)
130	c cgg gag aca cag atc tG (SEQ ID No:131)
131	g gct cag atc acc cag cA (SEQ ID No:132)
132	acc tac ctg gag ggc aC (SEQ ID No:133)
133	cac tcc atg agg tat ttc C (SEQ ID No:134)
134	gac ccc cca aag aca caT (SEQ ID No:135)
135	gag aca cag atc tcc aag aT (SEQ ID No:136)
136	gg gag gcg gcc cgt C (SEQ ID No:137)
137	gcg ccg tgg ata gag caA (SEQ ID No:138)
138	g acc aac aca cag act tac A (SEQ ID No:139)
139	ac acc ctc cag aat atg taT (SEQ ID No:140)
140	g gag ccc cgc ttc att G (SEQ ID No:141)
141	g gat tac atc gcc ctg aaG (SEQ ID No:142)
142	c acc ctc cag agg atg tG (SEQ ID No:143)
143	gcg ccg tgg ata gag caA (SEQ ID No:144)
144	cga gag aac ctg cgc aC (SEQ ID No:145)
145	gag aac ctg cgc acc gC (SEQ ID No:146)
146	g tct cac acc ctc cag aaT (SEQ ID No:147)
147	cag gag ggg ccg gag C (SEQ ID No:148)
148	ctg ggc ttc tac cct gG (SEQ ID No:149)
149	ca cag act gac cga gag G (SEQ ID No:150)
150	c gcc gcg gac acg gcA (SEQ ID No:151)

Table 5-6

Probe No.	Base Sequence
151	ctg ctc tgg ggg gca G ( SEQ ID No :152)
152	c cag agc gag gcc ggT ( SEQ ID No :153)
153	c tcc gtg tcc cgg ccT ( SEQ ID No :154)
154	cgc ggg tac cac cag C ( SEQ ID No :155)
155	tg acc gag acc tgg gcT ( SEQ ID No :156)
156	cag gag ggg ccg gag tT ( SEQ ID No :157)
157	cga gag agc ctg cgg aC ( SEQ ID No :158)
158	c acg gcg gct cag atc T ( SEQ ID No :159)
159	cg gag cag ctg aga gcT ( SEQ ID No :160)
160	gg ccc gac ggg cgc T ( SEQ ID No :161)
161	cgc ggg cat gac cag tT ( SEQ ID No :162)
162	cc atg tcc cgg ccc gT ( SEQ ID No :163)
163	g acc gcg gcg gac acC ( SEQ ID No :164)
164	c tgc gac gtg ggg ccC ( SEQ ID No :165)
165	t ccg agg acg gag ccC ( SEQ ID No :166)
166	gag ccc cgg gcg ccA ( SEQ ID No :167)
167	cc gcg agt ccg agg aC ( SEQ ID No :168)
168	cac atc atc cag agg atg tT ( SEQ ID No :169)
169	ca cag act tac cga gag. aA ( SEQ ID No :170)
170	c atg tac ggc tgc gac C ( SEQ ID No :171)
171	ctg cgg aac ctg cgc gA ( SEQ ID No :172)
172	cat gac cag tcc gcc tG ( SEQ ID No :173)
173	c acc atc cag agg atg tC ( SEQ ID No :174)
174	gac ctg agc tcc tgg acA ( SEQ ID No :175)
175	cga gag agc ctg cgc aC ( SEQ ID No :176)
176	g cag gag ggg ccg gG ( SEQ ID No :177)
177	ga acc tac ctg gag ggc A ( SEQ ID No :178)
178	a acc tac ctg gag ggc aT ( SEQ ID No :179)
179	c tgg acc gcg gcg gaG ( SEQ ID No :180)
180	ta gag cag gag ggg ccA ( SEQ ID No :181)

Table 5-7

Probe No.	Base Sequence
181	tct cac act tgg cag acG ( SEQ ID No :182)
182	g gcg gag cag cgg aga A ( SEQ ID No :183)
183	cgg ccc ggc cgc ggA ( SEQ ID No :184)
184	gg tct cac acc ctc caC ( SEQ ID No :185)
185	c cgc ggg tat aac cag ttA ( SEQ ID No :186)
186	g gcg gag cag tgg aga A ( SEQ ID No :187)
187	gaa tat tgg gac cgg gaG ( SEQ ID No :188)
188	gcg gct cag atc acc cG ( SEQ ID No :189)
189	cac acc ctc cag agc aC ( SEQ ID No :190)
190	ag tgg gag gcg gcc cT ( SEQ ID No :191)
191	g acc gag acc tgg gcG ( SEQ ID No :192)
192	c gcc acg agt ccg agg A ( SEQ ID No :193)
193	g atc tcc cag cgc aag tI ( SEQ ID No :194)
194	tg gag gcg gcc cgt gI ( SEQ ID No :195)
195	tg acc gag acc tgg gcI ( SEQ ID No :196)
196	g cgc tcc tgg acc gcG ( SEQ ID No :197)
197	ag ggc gag tgc gtg gaI ( SEQ ID No :198)
198	gg tat ttc cac acc gcc A ( SEQ ID No :199)
199	c cgc ggg cat aac cag A ( SEQ ID No :200)
200	ccg gag tat tgg gac cC ( SEQ ID No :201)
201	gg tct cac atc atc cag G ( SEQ ID No :202)
202	c gcc tac gac ggc aag A ( SEQ ID No :203)
203	cgc ggg cat aac cag tC ( SEQ ID No :204)
204	cc ggg tct cac act. tgG ( SEQ ID No :205)
205	c act tgg cag agg atg taI ( SEQ ID No :206)
206	ga gag agc ctg cgg aaG ( SEQ ID No :207)
207	c ggg aag gac acg ctg C ( SEQ ID No :208)
208	c acg ctg cag cgc gcG ( SEQ ID No :209)
209	cc atc tct gac cat gag gI ( SEQ ID No :210)
210	cgg gag aca cag atc tcG ( SEQ ID No :211)

Table 5-8

Probe No.	Base Sequence
211	g gag gcg gcc cgt gtC (SEQ ID No :212)
212	a gag aac ctg cgc acc G (SEQ ID No :213)
213	gg gag ccc cgc ttc atT (SEQ ID No :214)
214	ctg cgc acc ccg ctc C (SEQ ID No :215)
215	gg ccg gag tat tgg gaG (SEQ ID No :216)
216	c cgc ggg cat aac cag G (SEQ ID No :217)
217	ggc gag tgc gtg gag tC (SEQ ID No :218)
218	cgg gcg ccg tgg gtG (SEQ ID No :219)
219	ga gag aac ctg cgg atc G (SEQ ID No :220)
220	gtg gac gac acg ctg ttG (SEQ ID No :221)
221	tg gag ggc ctg tgc gC (SEQ ID No :222)
222	gac ggc aag gat tac atc A (SEQ ID No :223)
223	c cgc ggg tat aac cag tT (SEQ ID No :224)
224	ctc cgc ggg tat aac cG (SEQ ID No :225)
225	gcg gag cag gac aga gT (SEQ ID No :226)
226	gag aca cag aag tac aag C (SEQ ID No :227)
227	cgc cag gca cag act gG (SEQ ID No :228)
228	t gtg gtc gct gct gtg G (SEQ ID No :229)
229	c ctg cgg aac ctg ctc C (SEQ ID No :230)
230	aga acc ttc cag aag tgg A (SEQ ID No :231)
231	ag ccc cgc ttc atc tcC (SEQ ID No :232)
232	c cgc ggg tat aac cag ttA (SEQ ID No :233)
233	ggc ctg tgc gtg gag G (SEQ ID No :234)
234	cgg atc gcg ctc cgc G (SEQ ID No :235)
235	ttc gcc tac gac ggc aaA (SEQ ID No :236)
236	ctc ctc cgc ggg cat aaA (SEQ ID No :237)
237	g cgt ctc ctc cgc ggT (SEQ ID No :238)
238	c ggg cgc ctc ctc cC (SEQ ID No :239)
239	g agt ccg agg acg gag A (SEQ ID No :240)
240	ata gag cag gag ggg cG (SEQ ID No :241)

Table 5-9

Probe No.	Base Sequence
241	cc aga cca gca gga gat G (SEQ ID No :242)
242	cag cat gag ggg ctg cT (SEQ ID No :243)
243	cag act tac cga gag aac T (SEQ ID No :244)
244	gc gac gcc gcg agt cA (SEQ ID No :245)
245	c cgc ggg gag ccc cC (SEQ ID No :246)
246	cga gag agc ctg cgg aT (SEQ ID No :247)
247	gag agc ctg cgg atc gC (SEQ ID No :248)
248	g gca cag act gac cga gT (SEQ ID No :249)
249	g acc gcc gcg gac acC (SEQ ID No :250)
250	g cag gag ggg ccg gC (SEQ ID No :251)
251	cc gcg agt ccg aga gG (SEQ ID No :252)
252	gg tct cac act tgg cag aT (SEQ ID No :253)
253	acg gca ccc cga acc C (SEQ ID No :254)
254	ctc ctc ctg ctg ctc tG (SEQ ID No :255)
255	ag aca cag aag tac aag gG (SEQ ID No :256)
256	gg tct cac atc atc cag gT (SEQ ID No :257)
257	gc ggg cat gac cag tcT (SEQ ID No :258)
258	g acc gcg gcg gac acA (SEQ ID No :259)
259	g ccg gag tat tgg gac G (SEQ ID No :260)
260	c ctc ctc cgc ggg tat A (SEQ ID No :261)
261	c acg gcg gct cag atc aT (SEQ ID No :262)
262	tg cgg atc gcg ctc cC (SEQ ID No :263)
263	g ccg gag tat tgg gac gA (SEQ ID No :264)
264	g gag gcg gcc cgt gC (SEQ ID No :265)
265	c gac gcc gcg agt ccA (SEQ ID No :266)
266	gtc acc gta gct gtg gtC (SEQ ID No :267)
267	g tgt agg agg aag agt tcT (SEQ ID No :268)
268	c aga gcc tac ctg gag gA (SEQ ID No :269)
269	gtc atc gga gct gtg gtT (SEQ ID No :270)

Table 6-1

Probe No.	Base Sequence
0	c acc tcc Gtg tcc cgg (SEQ ID No:271)
1	c ctc cag agC atg tac gg (SEQ ID No:272)
2	c cgc ggg Cat gac cag (SEQ ID No:273)
3	cat gac cag Tac gcc tac (SEQ ID No:274)
4	g gag cag cGg aga gcc (SEQ ID No:275)
5	gag cag cGg aga gcc ta (SEQ ID No:276)
6	g gag ggc gAg tgc gtg (SEQ ID No:277)
7	c gtg gag tGg ctc cgc (SEQ ID No:278)
8	ac aag ctg Gag cgc gct (SEQ ID No:279)
9	ctc cgc agG tac ctg ga (SEQ ID No:280)
10	g gac gac acG cag ttc gt (SEQ ID No:281)
11	aag acc aac Aca cag act g (SEQ ID No:282)
12	g gag cag gaC aga gcc ta (SEQ ID No:283)
13	cgc ggg cat Aac cag tac (SEQ ID No:284)
14	cag tcc acc Atc ccc atc (SEQ ID No:285)
15	c ctc cag agG atg tac gg (SEQ ID No:286)
16	aca cag atc tTc aag acc aa (SEQ ID No:287)
17	t gac cag tCc gcc tac g (SEQ ID No:288)
18	ca cag atc tGc aag gcc c (SEQ ID No:289)
19	c cga gag aAc ctg cgg a (SEQ ID No:290)
20	tct cac atc Atc cag agg a (SEQ ID No:291)
21	g agg atg taT ggc tgc ga (SEQ ID No:292)
22	c tgc gac Ctg ggg ccc (SEQ ID No:293)
23	ctg ggg ccC gac ggg (SEQ ID No:294)
24	g tac aag cGc cag gca c (SEQ ID No:295)
25	ag gca cag Gct gac cga (SEQ ID No:296)
26	t gac cga gTg agc ctg c (SEQ ID No:297)
27	gg tct cac aTc atc cag ag (SEQ ID No:298)
28	c atc cag agG atg tac gg (SEQ ID No:299)
29	tc cgc ggg Tat gac cag (SEQ ID No:300)
30	aag acc aac Aca cag act ta (SEQ ID No:301)

Table 6-2

Probe No.	Base Sequence
31	aca cag act Tac cga gag a ( SEQ ID No :302)
32	g gag ggc aCg tgc gtg ( SEQ ID No :303)
33	ggg aag gaG acg ctg ga ( SEQ ID No :304)
34	g aag gag aCg ctg gag c ( SEQ ID No :305)
35	g gag ggc cTg tgc gtg ( SEQ ID No :306)
36	c gtg gag tCg ctc cgc ( SEQ ID No :307)
37	c ggg gag cTc cgc ttc ( SEQ ID No :308)
38	c gcc gcg Aac acg gcg ( SEQ ID No :309)
39	tg cgc ggc Gac tac aac ( SEQ ID No :310)
40	g gag ggc Ctg tgc gtg ( SEQ ID No :311)
41	g gcc cgt gTg gcg gag ( SEQ ID No :312)
42	g gag cag cTg aga gcc t ( SEQ ID No :313)
43	ca cag atc tCc aag acc aa ( SEQ ID No :314)
44	aca cag act Tac cga gag g ( SEQ ID No :315)
45	c cga gag Gac ctg cgg ( SEQ ID No :316)
46	cc ctg ctc Cgc tac tac ( SEQ ID No :317)
47	tat gac cag Gac gcc tac ( SEQ ID No :318)
48	agg tat ttc Gac acc gcc ( SEQ ID No :319)
49	c acc gcc Atg tcc cgg ( SEQ ID No :320)
50	gag ccg cCg gcg ccg ( SEQ ID No :321)
51	g gag ggc Acg tgc gtg ( SEQ ID No :322)
52	g agg aag agC tca ggt gg ( SEQ ID No :323)
53	cc gcg ctc Cgc tac tac ( SEQ ID No :324)
54	c ctg cgg aTc gcg ctc ( SEQ ID No :325)
55	g cgg atc gCg ctc cgc ( SEQ ID No :326)
56	tc gcg ctc Cgc tac tac ( SEQ ID No :327)
57	g aag gac aCg ctg gag c ( SEQ ID No :328)
58	ac aca cag aCc ttc aag ac ( SEQ ID No :329)
59	g acg atg taT ggc tgc ga ( SEQ ID No :330)
60	gg gac cgg Gac aca cag ( SEQ ID No :331)
61	ac cac cag Gac gcc tac ( SEQ ID No :332)



Table 6-3

Probe No.	Base Sequence
62	aac aca cag Gct gac cga (SEQ ID No:333)
63	gcc ctg ggC ttc tac cc (SEQ ID No:334)
64	c acc cag cTc aag tgg g (SEQ ID No:335)
65	ct tgg cag aCg atg tat gg (SEQ ID No:336)
66	t aac cag tTA gcc tac gac (SEQ ID No:337)
67	c tgc gac Ctg ggg ccg (SEQ ID No:338)
68	a tct tcc caA tcc acc gtc (SEQ ID No:339)
69	g aga gcc tGc ctg gag g (SEQ ID No:340)
70	acc ctc cag Tgg atg tat g (SEQ ID No:341)
71	a gca gga gaC aga acc ttc (SEQ ID No:342)
72	a tgg gag ccA tct tcc ca (SEQ ID No:343)
73	tc tac acc Gcc gtg tcc (SEQ ID No:344)
74	tcc atg agg Cat ttc tac ac (SEQ ID No:345)
75	g ggg ccg gaA tat tgg ga (SEQ ID No:346)
76	tc cgc aga Cac ctg gag (SEQ ID No:347)
77	g acg ctg Cag cgc gcg (SEQ ID No:348)
78	ctc tcg ggA gcc ctg g (SEQ ID No:349)
79	cgg gcg ccA tgg ata ga (SEQ ID No:350)
80	g gac cgg gaG aca cag at (SEQ ID No:351)
81	cg gag cag Tgg aga gcc (SEQ ID No:352)
82	t cag gac acC gag ctt gt (SEQ ID No:353)
83	c gac ggc aaA gat tac atc (SEQ ID No:354)
84	tgg acc gcG gcg gac a (SEQ ID No:355)
85	c gcc ctg aaI gag gac ct (SEQ ID No:356)
86	cag ttc gtg Cgg ttc gac (SEQ ID No:357)
87	gtg gtc gct Act gtg atg (SEQ ID No:358)
88	ag agg atg tTt ggc tgc g (SEQ ID No:359)
89	ca cag atc tGc aag acc aa (SEQ ID No:360)
90	agg atg gcI ccc cgg g (SEQ ID No:361)
91	tgc gtg gaC ggg ctc c (SEQ ID No:362)
92	gc tcc cac tTc atg agg t (SEQ ID No:363)

Table 6-4

Probe No.	Base Sequence
93	gcc tcc gcG cag act ta ( SEQ ID No :364)
94	tg gtg gtg cTt tct gga g ( SEQ ID No :365)
95	ac cac ccc Gtc tct gac ( SEQ ID No :366)
96	ac cgg gag aTa cag atc tc ( SEQ ID No :367)
97	g agg atg gGg ccc cgg ( SEQ ID No :368)
98	g agg atg tCt ggc tgc g ( SEQ ID No :369)
99	c gcg gac aAg gcg gct ( SEQ ID No :370)
100	cc ctc cag aGg atg tac g ( SEQ ID No :371)
101	c ctc cag acG atg tac gg ( SEQ ID No :372)
102	aac ctg cgC acc gcg c ( SEQ ID No :373)
103	ag gac ctg Agc tcc tgg ( SEQ ID No :374)
104	gc ttc atc Gca gtg ggc ( SEQ ID No :375)
105	atg gcg ccC cgg gcg ( SEQ ID No :376)
106	c gac gcc Acg agt ccg ( SEQ ID No :377)
107	cag ctg aga Acc tac ctg ( SEQ ID No :378)
108	cc aac aca cGg act tac c ( SEQ ID No :379)
109	ggg aAg gaG acg ctg ca ( SEQ ID No :380)
110	ac gac acg cTg ttc gtg a ( SEQ ID No :381)
111	ct tac cga gTg aac ctg c ( SEQ ID No :382)
112	c cga gtg aAc ctg cgg a ( SEQ ID No :383)
113	at aac cag tTc gcc tac ga ( SEQ ID No :384)
114	gtg agg ttc Aac agc gac ( SEQ ID No :385)
115	c acc cag cAc aag tgg g ( SEQ ID No :386)
116	cg gag cag cTg aga acc t ( SEQ ID No :387)
117	agg tat ttc C ac acc tcc g ( SEQ ID No :388)
118	a aag aca caT gtg acc cac ( SEQ ID No :389)
119	atc tcc aag aTc aac aca ca ( SEQ ID No :390)
120	g gcc cgt Cag gcg gag ( SEQ ID No :391)
121	g ata gag caA gag ggg cc ( SEQ ID No :392)
122	cag act tac Aga gag agc c ( SEQ ID No :393)
123	g aat atg taT ggc tgc gac ( SEQ ID No :394)

Table 6-5

Probe No.	Base Sequence
124	cgc ttc att Gca gtg ggc ( SEQ ID No :395)
125	gcc ctg aaG gag gac ct ( SEQ ID No :396)
126	ct tac cga gfg agc ctg c ( SEQ ID No :397)
127	g agg atg tGc ggc tgc g ( SEQ ID No :398)
128	g ata gag caA gag ggg cc ( SEQ ID No :399)
129	ca cag atc tGc aag gcc a ( SEQ ID No :400)
130	c ctg cgc aCc gcg ctc ( SEQ ID No :401)
131	cgc acc gCg ctc cgc ( SEQ ID No :402)
132	c ctc cag aaT atg tat ggc ( SEQ ID No :403)
133	gg ccg gag Cat tgg gac ( SEQ ID No :404)
134	tc tac cct gGg gag atc a ( SEQ ID No :405)
135	g gac acg gcA gct cag at ( SEQ ID No :406)
136	g ggg gca Gtg gcc ctg ( SEQ ID No :407)
137	gag gcc ggT tct cac ac ( SEQ ID No :408)
138	tcc cgg ccT ggc cgc ( SEQ ID No :409)
139	ac cac cag Cac gcc tac ( SEQ ID No :410)
140	acc tgg gcT ggc tcc c ( SEQ ID No :411)
141	g gtc acg gAg ccc cga ( SEQ ID No :412)
142	g ccg gag tTt tgg gac c ( SEQ ID No :413)
143	c ctc cag aaT atg tac ggc ( SEQ ID No :414)
144	c ctg cgg aCc ctg ctc ( SEQ ID No :415)
145	ct cag atc Tcc cag cgc ( SEQ ID No :416)
146	g ctg aga gcT tac ctg ga ( SEQ ID No :417)
147	c ggg cgc Ttc ctc cgc ( SEQ ID No :418)
148	at gac cag tTc gcc tac g ( SEQ ID No :419)
149	cgc ggg cat Aac cag ttc ( SEQ ID No :420)
150	cgg ccc gTc cgc ggg ( SEQ ID No :421)
151	gcg gac acC gcg gct c ( SEQ ID No :422)
152	tct cac atc Atc cag agc a ( SEQ ID No :423)
153	gtg ggg ccC gac ggg ( SEQ ID No :424)
154	acg gag ccC cgg gcg ( SEQ ID No :425)

Table 6-6

Probe No.	Base Sequence
155	t ccg agg aCg gag ccc ( SEQ ID No :426)
156	ac ctg cgc gAc tac tac a ( SEQ ID No :427)
157	g tcc gcc tGc gac ggc ( SEQ ID No :428)
158	tcc tgg acA gcg gcg g ( SEQ ID No :429)
159	c cga gag aAc ctg cgc a ( SEQ ID No :430)
160	g ggg ccg gGa tat tgg g ( SEQ ID No :431)
161	tg gag ggc Atg tgc gtg ( SEQ ID No :432)
162	g gag ggc aIg tgc gtg g ( SEQ ID No :433)
163	gcg gcg gaG acc gcg ( SEQ ID No :434)
164	g gag ggg ccA gaa tat tg ( SEQ ID No :435)
165	ct tgg cag aCg atg tac g ( SEQ ID No :436)
166	t tgg cag acG atg tac gg ( SEQ ID No :437)
167	cag cgg aga Acc tac ctg ( SEQ ID No :438)
168	ggc cgc ggA gag ccc ( SEQ ID No :439)
169	c acc ctc caC agg atg ta ( SEQ ID No :440)
170	cg gag cag Tgg aga acc ( SEQ ID No :441)
171	cag tgg aga Acc tac ctg ( SEQ ID No :442)
172	g atc acc cGg cgc aag t ( SEQ ID No :443)
173	c cag agc aCg tac ggc t ( SEQ ID No :444)
174	g gcg gcc cIt gtg gcg ( SEQ ID No :445)
175	acc tgg gcG ggc tcc c ( SEQ ID No :446)
176	gtc acg gcA ccc cga ac ( SEQ ID No :447)
177	agg tat ttc Cac acc gcc ( SEQ ID No :448)
178	gt ccg agg Aag gag ccg ( SEQ ID No :449)
179	g cgc aag tTg gag gcg g ( SEQ ID No :450)
180	acc tgg gcI ggc tcc c ( SEQ ID No :451)
181	tgc gtg gaT tgg ctc cg ( SEQ ID No :452)
182	cat aac cag Aac gcc tac g ( SEQ ID No :453)
183	t tgg gac cCg gag aca c ( SEQ ID No :454)
184	atc atc cag Gtg atg tat gg ( SEQ ID No :455)
185	gac ggc aag Aat tac atc g ( SEQ ID No :456)

Table 6-7

Probe No.	Base Sequence
186	at aac cag tCc gcc tac g (SEQ ID No :457)
187	ctg cgg aaG ctg cgc g (SEQ ID No :458)
188	t cac act tgG cag agg atg (SEQ ID No :459)
189	c acg ctg Cag cgc gcg (SEQ ID No :460)
190	ac cat gag gTc acc ctg a (SEQ ID No :461)
191	a cag atc tCG aag acc aac (SEQ ID No :462)
192	gcc cgt gtC gcg gag c (SEQ ID No :463)
193	g cgc acc Gcg ctc cg (SEQ ID No :464)
194	c cgc ttc atI gca gtg gg (SEQ ID No :465)
195	c ctg cgc aCc ccg ctc (SEQ ID No :466)
196	cc ccg ctc Cgc tac tac (SEQ ID No :467)
197	g tat tgg gaG cgg gag ac (SEQ ID No :468)
198	gc ggg cat Aac cag gac (SEQ ID No :469)
199	cat aac cag Gac gcc tac (SEQ ID No :470)
200	ctc cgc ggg Tat aac cag (SEQ ID No :471)
201	ccg tgg gtG gag cag g (SEQ ID No :472)
202	g cgg atc Gcg ctc cgc (SEQ ID No :473)
203	c acg ctg ttG gtg agg tt (SEQ ID No :474)
204	c ctg tgc gCg gag tcg (SEQ ID No :475)
205	gat tac atc Acc ctg aac g (SEQ ID No :476)
206	gg tat aac cGg tta gcc ta (SEQ ID No :477)
207	ag gac aga gTc tac ctg g (SEQ ID No :478)
208	aag tac aag Cgc cag gca (SEQ ID No :479)
209	ca cag act gGc cga gtg a (SEQ ID No :480)
210	gct gct gtg Gtg tgt agg (SEQ ID No :481)
211	aac ctg ctc Cgc tac tac (SEQ ID No :482)
212	cag aag tgg Aca gct gtg (SEQ ID No :483)
213	cag cgc gcG gac ccc (SEQ ID No :484)
214	c ttc atc tCt gtg ggc ta (SEQ ID No :485)
215	c gtg gag Ggg ctc cgc (SEQ ID No :486)
216	cg ctc cgc Gac tac aac (SEQ ID No :487)

Table 6-8

Probe No.	Base Sequence
217	c ggg cat aaA cag tac gc ( SEQ ID No:488)
218	c ctc cgc ggT tat aac ca ( SEQ ID No:489)
219	c ctc ctc cCc ggg cat ( SEQ ID No:490)
220	g acg gag Acc cgg gcg ( SEQ ID No:491)
221	g gag ggg cGg gag tat t ( SEQ ID No:492)
222	gca gga gat Gga acc ttc ( SEQ ID No:493)
223	g ggg ctg cTg aag ccc ( SEQ ID No:494)
224	cgg gtc aCg gcg ccc ( SEQ ID No:495)
225	t ccg agg aCg gag ccg ( SEQ ID No:496)
226	cga gag aac Ttg cgg atc ( SEQ ID No:497)
227	c gcg agt cAg agg acg g ( SEQ ID No:498)
228	g gag ccc cCc ttc atc g ( SEQ ID No:499)
229	g ggg ccg gCg tat tgg ( SEQ ID No:500)
230	t ccg aga gGg gag ccg ( SEQ ID No:501)
231	ct tgg cag aTg atg tat gg ( SEQ ID No:502)
232	g tac aag gGc cag gca c ( SEQ ID No:503)
233	tc atc cag gTg atg tat gg ( SEQ ID No:504)
234	t gac cag tcT gcc tac ga ( SEQ ID No:505)
235	gcg gac acA gcg gct c ( SEQ ID No:506)
236	tat tgg gac Ggg gag aca ( SEQ ID No:507)
237	cgc ggg tat Aac cag tac ( SEQ ID No:508)
238	ct cag atc aTc cag cgc a ( SEQ ID No:509)
239	c gcg ctç cCc tac tac a ( SEQ ID No:510)
240	at tgg gac gAg gag aca c ( SEQ ID No:511)
241	gcc cgt gCg gcg gag ( SEQ ID No:512)
242	g aag gag aCg ctg cag c ( SEQ ID No:513)
243	gcg agt ccA aga ggg ga ( SEQ ID No:514)
244	gct gtg gtC gct gtg gt ( SEQ ID No:515)
245	c ctg gag gAc ctg tgc g ( SEQ ID No:516)
246	a gct gtg gtT gct act gtg ( SEQ ID No:517)

## Table 7

## Allele-Probe List 1

	B*070201	0	1	2	3	4	5	6	7	8
	B*070202	9								
5	B*070203	10								
	B*0703	11								
	B*0704	12								
	B*0705	13	14							
	B*0706	13								
10	B*0707	15								
	B*0708	16	17							
	B*0709	18								
	B*0710	19								
	B*0711	20	18							
15	B*0712	21	22	23	24					
	B*0713	25	26	27						
	B*0714	28	21	29	30					
	B*0715	31	27							
	B*0716	11	32							
20	B*0717	30	33							
	B*0718	28	22							
	B*0719	12	34	35	36					
	B*0720	37	38							
	B*0721	39								
25	B*0722	40								
	B*0723	41								
	B*0724	42								

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20 B\*400103 177 110 106 178 80 43 31 15 13 179 41  
B\*4002 140 43 31 13 41 42 52  
B\*4003 140 80 43 31 17 41 42 52  
B\*4004 140 67 13 41 42 52  
B\*4005 140 13 42 35 52  
25 B\*400601 140 65 59 13 41 42 52  
B\*4007 16 31 15 13 179 41  
B\*4008 140 16 31 13 41 42 52

- B\*4009 117 110 106 29 41 42 109 77  
B\*4010 104 106 178 79 80 43 31 15 13 179 41  
B\*4011 117 110 106 15 13 41 42 109 77  
B\*4012 78 10 79 80 43 31 15 13 179 41  
5 B\*4013 140 55 56 13 41 42  
B\*401401 177 110 106 13 151 41 42  
B\*401402 177 110 106 13 84 41 42  
B\*4015 181  
B\*4016 176 43 30 31 13  
10 B\*4018 117 110 106 43 41 42 109 77  
B\*4019 106 43 30 54 13 41 42 77  
B\*4020 140 80 43 31 15 17 41 42  
B\*4021 105 79 80 43 31 15 13 179 41  
B\*4023 176 177 110 106 178 79 80 43 31 15 13 145  
15 B\*4024 117 106 79 80 43 30 31 22 23 41 42 77  
B\*4025 177 110 106 178 79 43 31 15 13 179 41  
B\*4026 106 80 43 31 83 42 35  
B\*4027 182  
B\*4028 67 83 42 35 76  
20 B\*4029 183 13  
B\*4030 184 179 42  
B\*4031 177 110 106 178 79 80 43 31 179 41  
B\*4032 177 49 110 106 79 80 43 31  
B\*4033 177 110 106 178 79 80 43 31 15 179 41  
25 B\*4034 185  
B\*4035 117 110 106 13 103 41 42 109 77  
B\*4036 15 113 179 41

- B\*4037 117 110 106 43 19 13 41 42 109 77  
B\*4038 186 179 41  
B\*4039 140 13 42 32 63  
B\*4040 110 106 43 13 41 42 109 77  
5 B\*4042 29 179 41  
B\*4043 177 110 106 178 79 80 43 15 13 179 41  
B\*4044 110 106 43 31 65 59 41 42 77  
B\*4101 176 21 12 32 52  
B\*4102 176 12 32 52  
10 B\*4103 168 12  
B\*4104 23 41 12 51 32 57  
B\*4105 187  
B\*4106 188 12 32 189  
B\*4201 12 32 52  
15 B\*4202 117 13 41 12 51 32 57  
B\*4204 65 59 13 41 12 51 32 57  
B\*440201 106 53 12 35 190 52  
B\*440202 191 53 12 35 190  
B\*440203 192  
20 B\*440301 106 42 35 190 52  
B\*440302 106 21 42 35 190 52  
B\*4404 32 190 52  
B\*4405 80 130 193 53 20 29 12 35 36 77  
B\*4406 194 12 36  
25 B\*4407 21 42 35 190 52  
B\*4408 105 53 12 35 190 52  
B\*4409 106 12 35 190 52

- B\*4410 88 42 36
- B\*4411 49 106 79 80 195 196 20 29 47 12 35 36 77
- B\*4412 49 106 79 130 193 53 20 29 47 12 35 36 77
- B\*4413 197
- 5 B\*4414 198 199 12 35 36 77
- B\*4415 130 53 200 66 12 36
- B\*4416 49 106 79 80 102 193 20 29 47 12 77
- B\*4417 17 12 35 36 77
- B\*4418 176 56 200 12 36 77
- 10 B\*4420 59 29 47 12 35 36 77
- B\*4421 49 106 79 80 130 193 53 20 29 47 12 36 77
- B\*4422 106 79 80 130 193 53 20 29 47 12 35 36 77
- B\*4424 201 36 77
- B\*4425 43 54 202 56 20 29 12 35 36 77
- 15 B\*4426 203
- B\*4427 106 53 47 12 35 36 82
- B\*4428 106 20 21 29 47 151 41 35 36 77
- B\*4429 107 36 77
- B\*4430 204
- 20 B\*4431 179 36
- B\*4432 183 36
- B\*4433 205
- B\*4501 176 200 12 36 82 87 52
- B\*4502 31 200 113 12 35 36 77
- 25 B\*4503 206
- B\*4504 176 200 12 82 87 52
- B\*4505 207

- B\*4506 200 66 12 36  
B\*4601 208 52  
B\*4602 209  
B\*470101 210  
5 B\*4702 88 41 42 63  
B\*4703 211 88 61 41 42  
B\*4704 30 130 131 53 132 61 41 42 77  
B\*4801 31 212  
B\*4802 43 30 20 21 17 103 41 42 35 95 52  
10 B\*4803 80 43 31 15 13 179 41  
B\*4804 213 212  
B\*4805 214 80 43 31 13  
B\*4806 16 31 13 179 41  
B\*4807 113 179 41  
15 B\*4901 176 56 42 82 87 52  
B\*4902 130 53 66 42 82  
B\*4903 154 54 56 22 200 66 42 35  
B\*5001 176 42 82 87 52  
B\*5002 176 42 36 82 87 52  
20 B\*5004 106 200 66 42 35  
B\*510101 194 16 56 65 42 76 95 52  
B\*510102 16 56 65 42 76 95 52  
B\*510103 194 16 56 65 59 83 103 42 35 76  
B\*510104 194 79 16 56 65 59 13 103 42 35 76  
25 B\*510105 118 95 87  
B\*510201 194 16 56 65 83 103 42 35 95 52  
B\*510202 16 56 65 83 103 42 35 95 52

- B\*5103 215 95  
B\*5104 20 42 76 95 52  
B\*5105 56 83 103 41 35 95  
B\*5106 194 79 16 56 83 103 42 35 76  
5 B\*5107 194 155 43 56 65 59 83 103 42 35 76  
B\*5108 12 76 95 52  
B\*5109 194 41 42 35 76  
B\*5110 56 65 59 13 41 42 95  
B\*5112 216  
10 B\*511301 194 113 103 42 35 76  
B\*511302 194 113 83 103 42 35 76  
B\*5114 217  
B\*5115 54 56 65 59 67 66 41 42 35  
B\*5116 194 79 16 56 65 59 83 103 42 76  
15 B\*5117 218  
B\*5118 219  
B\*5119 194 107 35 76  
B\*5120 194 12 35 76  
B\*5121 194 83 32 76  
20 B\*5122 194 89 56 65 59 83 103 42 35 76  
B\*5123 194 42 36  
B\*5124 194 79 16 56 65 59 13 42 35 76  
B\*5126 220  
B\*5128 221  
25 B\*5129 194 16 56 65 76 95  
B\*5130 95 222  
B\*5131 194 83 41 42 76



- B\*5132 223
- B\*5133 92 76
- B\*5134 194 79 16 56 65 59 83 103 151 42
- B\*520101 224 43 56 65 42 76 95 52
- 5 B\*520102 194 43 56 65 42 76 95 52
- B\*520103 225 80 43 56 65 59 83 42 35 76
- B\*520104 226
- B\*5202 194 97 80 43 56 65 59 83 103 42 35 76
- B\*5203 194 43 41 42 35
- 10 B\*5204 227
- B\*5205 228
- B\*5301 30 56 20 21 17 103 41 42 35 95 52
- B\*5302 56 17 41 42 35 76
- B\*5303 45 42 35 77
- 15 B\*5304 54 56 23 148 103 151 41 42 35 77
- B\*5305 30 54 55 56 20 21 22 23 17 103 151 41 42 35 77
- B\*5306 194 17 103 42 35 76
- B\*5307 98 42 35
- B\*5308 30 19 54 55 56 20 21 22 23 17 103 151 42 35 77
- 20 B\*5309 102 17 151 41 42 40 35 77
- B\*5401 86 52
- B\*5402 117 86 32
- B\*5501 176 32 87 52
- B\*5502 176 41 32 87 52
- 25 B\*5503 26 66 32 77
- B\*5504 49 104 10 13 151 41 42 32 109 77
- B\*5505 229

- B\*5507 49 230 66 77
- B\*5508 104 10 15 13 151 41 42 35 77
- B\*5509 49 104 10 65 59 67 66 151
- B\*5510 65 59 67 66 41 32
- 5 B\*5511 231 32 77
- B\*5512 176 19 41 32 87 52
- B\*5601 176 59 41 42 35 87 52
- B\*5602 176 41 42 35 87 52
- B\*5603 176 81 82 87 52
- 10 B\*5604 104 10 66 41 42 35
- B\*5605 194 10 65 59 83 103 42 35 76
- B\*5606 194 155 79 65 59 83 103 42 35 76
- B\*5607 10 130 53 65 59 67 66 41 42 35
- B\*5608 232 42 35 77
- 15 B\*5609 104 10 20 21 22 23 17 103 151 41 42 35 77
- B\*5610 49 104 10 67 66 41 32
- B\*5611 176 23 151 41 42 35 77 63
- B\*570101 233 17 42 68
- B\*570102 234
- 20 B\*5702 13 68
- B\*570301 13 42 68
- B\*570302 235
- B\*5704 184 47 41 77
- B\*5705 236 200 237 41 35
- 25 B\*5706 238
- B\*5707 184 36 77
- B\*5708 239

- B\*5709 184 12 77  
B\*5801 236 20 87 52  
B\*5802 70 52  
B\*5804 240  
5 B\*5805 241  
B\*5806 70 35  
B\*5807 70 36  
B\*5901 176 56 41 32 87 52  
B\*670101 75 15 116 107 71 72  
10 B\*670102 15 149 113 41 116 107 32 242  
B\*6702 243  
B\*7301 244  
B\*7801 194 16 65 42 76 95 52  
B\*780201 16 31 65 42 76 95 52  
15 B\*780202 194 79 16 31 65 59 83 103 42 35 76  
B\*7803 194 89 11 65 59 83 103 42 35 76  
B\*7804 83 103 41 42 35 95  
B\*7805 155 154 80 43 31 65 59 83 42 35 76  
B\*8101 136 212  
20 B\*8201 245  
B\*8202 246  
B\*8301 136 49 20 29 47 12 35 36 77

(Example 5)

Probes for identification of HLA-C allele

Extraction of DNA from 1 ml of human blood was performed using GFX Genomic Blood DNA Purification

5 Kit from Amersham Biosciences in the same manner as in Example 1.

Next, quantitative PCR was carried out in the same manner as in Example 1 except that probes in the probe list in Tables 9-1 to 9-4 were used

10 respectively, and 3  $\mu$ l of the mixed primers consisting of 1  $\mu$ l each of the respective solutions of the following primers (10 pmol/ $\mu$ l) was used:

AAACACGGTCACCTCAGGGGAT (SEQ ID NO: 340)

GGCCTGAGTGTGGTTGGAACG (SEQ ID NO: 341)

15 CCAGCTCGTAGTTGTGTCTGCA (SEQ ID NO: 342).

After PCR amplification, the sample was identified being Cw\*120202, referring to Amp Plot and Dissociation curves on a display of 5700 software and the allele-probe list in Tables 11-1 to 11-4.

20 (Example 6)

Extraction of DNA from 1 ml of human blood was performed in the same manner as in Example 1. PCR of human HLA-C was then performed in the same manner as in Example 2 except that 6  $\mu$ l of the mixed primer

25 consisting of 1  $\mu$ l each of the solutions containing the following sequences at 10 pmol/ $\mu$ l respectively and 9  $\mu$ l of ultra pure water was used.

AAACACGGTCACCTCAGGGGGAT (SEQ ID NO: 340)

GGCCTGAGTGTGGTTGGAACG (SEQ ID NO: 341)

CCAGCTCGTAGTTGTGTCTGCA (SEQ ID NO: 342)

CCATGTGTCAACTTATGCC (SEQ ID NO: 343)

5 AGAATTACCTTTTCCAG (SEQ ID NO: 344)

AGAATTACGTTTTCCAG (SEQ ID NO: 345)

At the same time, a DNA microarray was prepared to identify the allele in the specimen in the same manner as in Example 2. Probes in Tables 10-1 to 10-4 were used for the probe spots respectively.

Then, hybridization and fluorescence determination was performed using the above-prepared sample and the DNA microarray in the same manner as in Example 2 and the sample was identified as Cw\*120202 referring to the probe-allele list in Tables 12-1 to 12-4.

#### Allele list

Cw\*0102 :

20 atgcgggtcatggcgccccgaacctcatccigtgtctctcgggagccctggccctgaccgagacctgggcctgct  
cccactccatgaaglatcttccacatccgtgtccggccctggcccgaggagagccccgccttcatctcagtgaggcta  
cgtggacgacacgcagttcgtgcggttcgacagcgacgccgagtcaggagaggggagccgcggcgccgtgggtg  
gagcaggagggggccggagtatgggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtgagcc  
tgcggaacctgcgcggttactacaaccagagcgaggccgggtctcacacctccagtggatgtGtggctgcgacct  
25 ggggccccgacggcgccctccctccgggtatgaccagtAcgcctacgacggcaaggattacatcgccctgaacgag  
gacctgcgtccttgaccgcccgggacacCcggttcagatcaccagcgcaagtgaggagggcccggtgaggcgg  
agcagcgagagacctaccitggagggcacgtgcgtggagtggctccgcagataccitggagaacgggaaggagacgt

gcagcgcgcggaacacccaaagacacacgtgacccaccaatcccgtctctgaccaatgaggccacccatgaggctgtgg  
gcccctgggcttcttacctgctggagatcacactgacctggcagtgggatggggaggaccaaacctcaggacaccgagc  
ttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtgAtggctgcttctggagaagagca  
gagatacacgtgccatgtgcagcacgaggggctgccggagccccctacccctgagatgggagccgcttctccagccc  
5 accatccccatcgtgggcatcgttgcctggcctggctgtctctaggctgtctctaggagctgtggctggctg  
ttgtgagtgttaggaggaagagctcaggctggaaaaggaggaggagctgtctctcaggctgcgtccagcaacagtgtccca  
gggctctgagtgtctctcatcgttgttaa (SEQ ID NO:1);

Cw\*0103 :

atgcgggtcatggcgccccgaacccctcatctgtctgtctctcgggagccccggccccgacctgaggacccctggcctgtct  
10 cccactccatgaagtaattcttccatccgtgtctccggccctggccgcggagagccccgcttcatctcagtgggctta  
cgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagctccgagaggggagccgcgggcgccgtgggtg  
gagcaggagggggccggagtaattgggaccgggagacacagaagtaacaagcgccaggcacagactgacctgagtgacc  
tgcggaacctgcgcggctactacaaccagagcgaggccgggtctcacaccttccagtggaatgtgtggctgcgacct  
ggggcccgacgggcgcctctctccgggtatAaccagttcgcctacgacggcaaggattacatcgcctgaacgag  
15 gacctgcgtctctggaccgcccggacaccgcggctcagatcacccagcgcaagtgggagggcgccccgtgaggcgg  
agcagcggagagcctacctggagggcacgtgcgtggagtggtctccgcagataacctggagaacgggaaggagacgt  
gcagcgcgcggaacacccaaagacacacgtgacccaccaatcccgtctctgaccaatgaggccacccatgaggctgtgg  
gcccctgggcttcttacctgctggagatcacactgacctggcagtgggatggggaggaccaaacctcaggacaccgagc  
ttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtgattggcttcttggagaagagca  
20 gagatacacgtgccatgtgcagcacgaggggctgccggagccccctacccctgagatgggagccgcttctccagccc  
accatccccatcgtgggcatcgttgcctggcctggctgtctctaggctgtctctaggagctgtggctggctg  
ttgtgagtgttaggaggaagagctcaggctggaaaaggaggaggagctgtctctcaggctgcgtccagcaacagtgtccca  
gggctctgagtgtctctcatcgttgttaa (SEQ ID NO:2);

Cw\*0104 :

25 atgcgggtcatggcgccccgaacccctcatctgtctgtctctcgggagccccggccccgacctgaggacccctggcctgtct  
cccactccatgaagtaattcttccatccgtgtctccggccctggccgcggagagccccgcttcatctcagtgggctta  
cgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagctccgagaggggagccgcgggcgccgtgggtg

gagcaggaggggccggagatattgggaccgggagacacagaagtlacaagcgccagggcacagactlgaccgagtlgagcc  
tgcggaacctlgcgcggctactacaaccagagcgaggccgggtctcacaccttccagltggatgtgtggctlgcgacct  
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gacctlgcgttctlggaccgcTgcggacacggcggctcagatcacccagcgcaagtgggaggcggcccgtagggcgg  
5 agcagltggagagcctacctggagggcacgtlgcgtggagltggcttccgcagataacctggagaacgggaaggagacgct  
gcagcgcgcggaacacccaaagacacacgtgaccacccatcccgtctctlgacatgaggccaccttaggtgtctgg  
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ttgtggagaccaggccagcaggagatggaaccttcagaagtgggcagctgttggtggltgccttctlggagaagagca  
gagatacacgtgccatgtgcagcacgaggggtgccAgagccccctacccctgagatgggagccatcttcccagccc  
10 accttccccatcgtgggcacgttgcgtggcctggctgttctggctgttcttagctgttctaggagctgtgAtggctg  
ttgtgatgtgtaggaggaagagctcaggtggaaaaggaggaggagctgtcttcaggctgcgtccagcaacagtgccca  
gggctcttgatgagctctctcatcgcttgttaa (SEQ ID NO:2);

CW\*0105 :

15 gctccaccatccatgaaatattcttcacatccgtgtcccgccctggccgcggagagccccgccttcacatcagttggg  
ctacgtggacgacacgcagttctgtgcggttcgacagcgacgccgcgagtcgagagggggagccgcgggcgccgtgg  
gtggagcaggagggggccggagtatigggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtga  
gccttgcggaaccttgcgcggctactacaaccagagcgaggccgggtctcacaccttcagaggatgtctggctgcga  
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gaggacctgcgctccctggaccgccgcggacaccgcggctcagatcaccagcgcaagctgggaggcggcccgctgagg  
20 cggagcagcggagagccctacctggagggcacgtgcgtggagttggctccgcagatacctggagaacgggaaggagac  
gctgcagcgcgcgg (SEQ ID NO:4);

Cw\*0106 :

gc tccac tcca tgaagt att ctt caca tccgt gtc ccg gcc tgg ccg cgg agag ccc cgt tca tct cagt ggg  
ct acgt gga cgac acgc agt tct gtc ggt tgc acag cgac gcc cgc agt ccg agag ggg agc cgc ggg cgc cgt ggg  
25 g tgg agc agg aggg ggc cgg agt att ggg acc ggg agac acaga agt aca agc gcc aggc acag ac t gacc gagt ga  
gcc tgc gga acc tgc gcg gct ac taca acc agag cgagg ccgg gct tca cacc ctc agt gga tgt gt ggc tgc ga  
cc tgg ggc ccg acg ggc gcc tct ccc ggg gta t gacc agt acgc ctac gac ggca agga t taca tgc cc tga ac

gaggaccigcgc tcc tggaccgccg cgacaccgcggc tca gacacccagcgca agtgggaggcgcccg tTgg  
cggagcagcggagagcc tacc tggagggcacgtgc tggagtggc tccgcagat acc tggagaacgggaaggagac  
gctgcagcgcgcgg (SEQ ID NO:5);

Cw\*0107:

5 gctcccac tcca tgaagtatt tcttcacatccgtgt cccggcc tggccgaggagccccgc ttcattcagtggg  
ctacgtggacgacacgc agt tgc tgcgg ttcgacagcgacgccgcg agtccgagaggggagccgcgggcgcctgg  
gtggagcaggaggggcccgg agtattgggaccgggagacacaga agtacaagcggcaggcacagac taccgagtga  
gcc tgcggaacctgcgcggctactaca accagagcgaggccgggtctcacacctcc agtggatgtgtggctgcga  
cc tggggcccgacgggcgcctcctccgcAggtatgacc agt acgcctacgacggcaaggattacatcgccctgaac  
10 gaggaccigcgc tcc tggaccgccg cgacaccgcggc tca gacacccagcgca agtgggaggcgcccg tTgagg  
cggagcagcggagagcc tacc tggagggcacgtgc tggagtggc tccgcagat acc tggagaacgggaaggagac  
gctgcagcgcgcgg (SEQ ID NO:6);

Cw\*0108:

gctcccac tcca tgaagtatt tcttcacatccgtgt cccggcc tggccgaggagccccgc ttcattcagtggg  
15 ctacgtggacgacacgc agt tgc tgcgg ttcgacagcgacgccgcg agtccgagaggggagccgcgggcgcctgg  
gtggagcaggaggggcccgg agtattgggaccgggagacacaga agtacaagcggcaggcacagac taccgagtga  
gcc tgcggaacctgcgcggctactaca accagagcgaggccgggtctcacacctcc agtggatgtgtggctgcga  
cc tggggcccgacgggcgcctcctccgcgggtatgacc agt acgcctacgacggcaaggattacatcgccctgaac  
gaggaccigcgc tcc tggaccgccg cgacaccgcggc tca gacacccagcgca agtgggaggcgccTgtgagg  
20 cggagcagcggagagcc tacc tggagggcacgtgc tggagtggc tccgcagat acc tggagaacgggaaggagac  
gctgcagcgcgcgg (SEQ ID NO:7);

Cw\*0109:

gctcccac tcca tgaagtatt tcttcacatccgtgt cccggcc tggccgaggagccccgc ttcattcagtggg  
ctacgtggacgacacgc agt tgc tgcgg ttcgacagcgacgccgcg agtccgagaggggagccgcgggcgcctgg  
25 gtggagcaggaggggcccgg agtattgggaccgggagacacaga agtacaagcggcaggcacagac taccgagtga  
gcc tgcggaacctgcgcggctactaca accagagcgaggccgggtctcacacctcc agtggatgtgtggctgcga  
cc tggggcccgacgggcgcctcctccgcgggtatgacc agt acgcctacgacggcaaggattacatcgccctgaac





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ttgtggagaccaggccagcaggagatggaaccttccagaatgggcagctgtggctggcttctggagaagagca  
5 gagatacacgtgccatgtgcagcacgagggcgctccggagccccctacccatgagatgggagccatcttccagccc  
accatccccatcgtgggcatcgttgcctggcctggctgtccctggctgtcctagctgtcctaggagctgtggctggctg  
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gggctctgatgagctctcactcgttgtaa (SEQ ID NO:10);

Cw\*020203 :

10 gctcccacitccatgaggtatttctacaccgctgtgtcccgcccagccgcggagagccccacttcatcgcagtggg  
ctacgtggacgacacgcagttcgtgcggctcgacagcgacccgcgagttcaagaggggagccgcggcgccgtgg  
gtggagcaggaggggcccggagtattgggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtga  
acctgcggaaactgcgcggctactacaaccagagcgaggccgggtctcacacctccagaggatgtatggctgcga  
ccctggggcccgacggcgccctctccgcgggtatgaccagtcgcctacgacggcaaggattacatgccccgaac  
15 gaggacctgcgtctctggaccgcccggacacAgcggctcagatcacccagcgcaagtgggaggcgccccgtgagg  
cggagcagTggagagccatccatggaggcgagtcgctggagtggctccgcagataccatggagaacgggaaggagac  
gctgcagcgcgcgg (SEQ ID NO:11);

Cw\*020204 :

atgcgggtcatggcgccccgaaccttctctctgtctgtctcgggagccccggccccgaccgagacctgggctgtct  
20 cccacitccatgaggtatttctacaccgctgtgtcccgcccAgccgcggagagccccacttcatcgcagtgggcta  
cgtggacgacacgcagttcgtgcggctcgacagcgacccgcgagttcaagaggggagccgcggcgccgtgggtg  
gagcaggaggggcccggagtattgggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtgaacc  
tgcggaaactgcgcggctactacaaccagagcgaggccgggtctcacacctccagaggatgtacggctgcgacct  
ggggcccgacggcgccctctccgcgggtatgaccagtcgcctacgacggcaaggattacatgccccgaacgag  
25 gacctgcgtctctggaccgcccggacacGgcggctcagatcacccagcgcaagtgggaggcgccccgtgaggcgg  
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gcagcgcgcggaacacccaaagacacacgtgaccaccatcccgctcttgaccatgaggccacccatgaggctgtgg

gccccgggcttctacccitgccgagatcacactgacctggcagcgggaiggcaggaccaaactcaggacaccgagc  
ttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtggagggtgcttctggagaagagca  
gagatacacgtgccatgtgcagcacgaggggctgccggagccccctacccctgagatgggagccAtcttccagccc  
accaiccccatcgtgggcaicgttgcctggccctggcgttccctggcgttccctagctgtcctaggagctgtggaggctg

5 ttgtgatgtgtaggaggaagagctcag (SEQ ID NO:12);

Cw\*020205 :

gtccccactccatgaggatattctacaccgctgtgtccccggcccAgccgcggagagccccacttcatcgcagtggg  
ctacgtggacgacacgcagttcgtgcgggttcgacagcgacgccgcgagltccaagaggggagccgcgggcccgtgg  
gtggagcaggagggggccggagtattgggaccgggagacacagaagtacaagcgccaggcacagacigaccgagtga  
10 acctgcggaaactgcgcggctactacaaccagagcgaggccgggtctcacacctccagAggatgtatggctgcga  
cctggggcccgacgggcccctccctccgcggtatgaccagttcgccctacgacggcaaggattacatcgccctgaac  
gaggacctgcgctccctggaccgccgaggacacGgcggctcagatcacccagcgcaagtgaggagcgggcccgtgagg  
cggagcagTggagagccctacctggagggcgAgctgcgtggagtggctccgcagatacctggagaacgggaaggagac  
gctgcagcgcgcgg (SEQ ID NO:13);

15 Cw\*0203 :

gtccccactccatgaggatattctacaccgctgtgtccccggcccagccgcggagagccccacttcatcgcagtggg  
ctacgtggacgacacgcagttcgtgcgggttcgacagcgacgccgcgagltccaagaggggagccgcgggcccgtgg  
gtggagcaggagggggccggagtattgggaccgggagacacagaagtacaagcgccaggcacagacigaccgagtga  
acctgcggaaactgcgcggctactacaaccagagcgaggccgggtctcacacctccagaggatgtacggctgcga  
20 cctggggcccgacgggcccctccctccgcggtatgaccagttcgccctacgacggcaaggattacatcgccctgaac  
gaggacctgcgctccctggaccgccgaggacacagcggctcagatcacccagcgcaagtgaggagcgggcccgtTgg  
cggagcagctgagagccctacctggagggcgAgctgcgtggagtggctccgcagatacctggagaacgggaaggagac  
gctgcagcgcgcgg (SEQ ID NO:14);

Cw\*0204 :

25 gtccccactccatgaggatGttctacaccgctgtgtccccggcccagccgcggagagccccacttcatcgcagtggg  
ctacgtggacgacacgcagttcgtgcgggttcgacagcgacgccgcgagltccaagaggggagccgcgggcccgtgg  
gtggagcaggagggggccggagtattgggaccgggagacacagaagtacaagcgccaggcacagacigaccgagtga

accatgcggaactgcgcggctactacaaccagagcgaggccgggtctcacacccctccagaggatgtacggctgcga  
ccatggggcccgacgggcgcctctccgcgggtatgaccagtcgcctacgacggcaaggattacatgccccgaac  
gaggacctgcgctcctggaccgccgaggacacagcggtcagatcacccagcgcaagtgaggaggcgcccgctgagg  
cggagcagtgagagcctaccatggaggcgagtcgtggagtggtccgcagataccatggagaacgggaaggagac  
5 gctgcagcgcgcg(SAQ ID NO:15);

Cw\*0205 :

gctcccatccatgaggatattctacaccgctgtgtcccgccccAgccgcggagagccccacttcatgcagtgagg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgaagaggggagccgcgggcgcctgg  
gtggagcaggagggcgaglatgggaccgggagacacagaaglacaaagccaggcacagactgaccgagtga  
10 accatgcggaactgcgcggctactacaaccagagcgaggccgggtctcacacccctccagtggaatgtatggctgcga  
ccatggggcccgacgggcgcctctccgcgggtatgaccagtcgcctacgacggcaaggattacatgccccgaac  
gaggacctgcgctcctggaccgccgaggacacGgcggctcagatcacccagcgcaagtgaggaggcgcccgctgagg  
cggagcagtgagagcctaccatggaggcgAgatgcgtggagtggtccgcagataccatggagaacgggaaggagac  
gctgcagcgcgcg(SAQ ID NO:16);

15 Cw\*0206 :

gctcccatccatgaggatattctacaccgctgtgtcccgccccagccgcggagagccccacttcatgcagtgagg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgaagaggggagccgcgggcgcctgg  
gtggagcaggagggcgaglatgggaccgggagacacagaaglacaaagccaggcacagactgaccgagtga  
accatgcggaactgcgcggctactacaaccagagcgaggccgggtctcacacccctccagaggatgtacggctgcga  
20 ccatggggcccgacgggcgcctctccgcgggtatgaccagttAgcctacgacggcaaggattacatgccccgaac  
gaggacctgcgctcctggaccgccgaggacacggcggtcagatcacccagcgcaagtgaggaggcgcccgctgagg  
cggagcagtgagagcctaccatggaggcgAgatgcgtggagtggtccgcagataccatggagaacgggaaggagac  
gctgcagcgcgcg(SAQ ID NO:17);

Cw\*030201 :

25 atgcgggtcatggcggccgaacccctcatcctgtgtctcgggagccctggccctgaccgagacctgggcccgt  
cccatccatgaggatattctacaccgctgtgtcccgccccggccgcgggagccccacttcatgcagtgaggct  
cgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagccgcgggcgcctgggtg

gagcaggagggcgaggatattgggaccgggagacacagaagtacaagcgccaggcacagacigaccgagtagacc  
tgcggaacctgcgcggctactacaaccagagcgaggccgggttcacatccctccagaggatgtatggctgcgacgt  
ggggcccgacgggcgcctccctccgcggtatgaccagtcgcctacgacggcaaggattacatcgccctgaacgag  
gatctgcgctccctggaccgcccgggacacggcggtcagatcaccagcgcaagtgggaggcgcccgtagggcgg  
5 agcagctgagagcctaccctggagggcctgtgcgtggagtggctccgcagatacctgaagaatgggaaggagacgct  
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gcccctgggcttctacccctgcggagatcacactgacctggcagtaggaatgggaggaccacaaacacaggacacTgagc  
ttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtggtaggtgcttctggagaagagca  
gagatcacgtgccatgtgcagcacgaggggctgccggagccccctacccctgagatgggagccAtcttcccagccc  
10 accatccccatcgtagggcatcgtagctggccctggctgtccctggctgtccctagctgtccctaggagctgtggtaggctg  
ttgtgatgtgtaggaggaagagctcaggtaggaaaaggaggagctgtctcaggctgcgtccagcaacagtagccca  
gggctctgatgagctctctcatcgcttcttaa (SEQ ID NO:18);

Cw\*030202 :

atcggggtcatggcgccccgaacctcatctgtctgtctcgggagccccggccccgaccgagacctggggcggt  
15 cccactccatgaggatattctacaccgtgtgtccggccccggcggggagccccacttcatcgagtagggcta  
cgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagctccgagaggggagcccgcgggcgccgtgggtg  
gagcaggagggcgaggatattgggaccgggagacacagaagtacaagcgccaggcacagacigaccgagtagacc  
tgcggaacctgcgcggctactacaaccagagcgaggccgggttcacatccctccagaggatgtatggctgcgacgt  
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20 gatctgcgctccctggaccgcccgggacacggcggtcagatcaccagcgcaagtgggaggcgcccgtagggcgg  
agcagctgagagcctaccctggagggcctgtgcgtggagtggctccgcagatacctgaagaatgggaaggagacgct  
gcagcgcgcggaacacccaaagacacacgtgaccaccatcccgtctctgaccatgaggccaccctgaggtgctgg  
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ttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtggtaggtgcttctggagaagagca  
25 gagatcacgtgccatgtgcagcacgaggggctgccggagccccctacccctgagatgggagccgtcttcccagccc  
accatccccatcgtagggcatcgtagctggccctggctgtccctggctgtccctagctgtccctaggagctgtggtaggctg  
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gggcctcgtgagtcctcctcgttgtaa (SEQ ID NO:19);

Cw\*030301:

atgcgggtcatggcggccgaacccatcctgctgctcctgggagcccggcccagaccagaccggccggct  
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5 cgtggacgacacgcagtcgtgcggttcgacagcgacccgcgagtcgagaggggagccggggcgccgtgggtg  
gagcaggagggggcgagtatgggaccgggagacacagaagtacaagcgccaggcacagactgaccagtgagcc  
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10 agcagctgagagcctacctggagggcctgtgctggagtggtccgcagatacctgaagaatgggaaggagacgt  
gcagcgcgcggaacacccaaagacacacgtgaccacatcccgtctctgaccatgaggccacctgaggtgctgg  
gcccgggctcttacctgaggagatcacactgacctggcagtggtatggggaggaccaaactcaggacactgagc  
ttgtggagaccaggccagcaggagatggaaccttcagaagtgggcagctgtGtggtgctctctggagaagagca  
gagatcacgtgccatgtgcagcagcaggggctgccggagccccacccagatgggagccgtcttccagccc  
15 acctccccatcgtgggcatcgtgtgctggcctggctgtctggctgctctagctgtctctaggagctgtgggtggctg  
ttgtgatgtgtaggaggaagagctcaggtggaaaaggaggagctgctctcaggctgcgtccagcaacagtgccca  
gggcctcgtgagtcctcctcgttgtaa (SEQ ID NO:20);

Cw\*030302:

atgcgggtcatggcggccgaacccatcctgctgctcctgggagcccggcccagaccagaccggccggct  
20 cccacccaatgaggtatctcaccgcctgctccggccggccggggagccccacttcacgcagtgggcta  
cgtggacgacacgcagtcgtgcggttcgacagcgacccgcgagtcgagaggggagccggggcgccgtgggtg  
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25 gatctgcgtctctggaccggcgggacacggcggtcagatcaccagcgcaagtgaggagggcgccgtgaggcgg  
agcagctgagagcctacctggagggcctgtgctggagtggtccgcagatacctgaagaatgggaaggagacgt  
gcagcgcgcggaacacccaaagacacacgtgaccacatcccgtctctgaccatgaggccacctgaggtgctgg

gccc tgggct tctaccc tgcggagatcacactgacctggcag tggga tggggaggaccaaactcaggacac t gaggc  
t t g tggagaccaggccagcaggagatggaacct tccagaag tgggcagc t g tgg tgg tgcct tctggagaagagca  
gagatacacgtgccatgtgcagcacgaggggctgccggagcccc taccctgagatgggagccg tct tccagccc  
accatccccatcgtgggcatcgt tgc tggcctggc t g cctggc t g cctagc t g cctaggagc t g tgg tggc t g  
5 t t g t g a t g t g t a g g a g a a g a g c t c a g g t g g a a a g g a g g g a g c t g c t c t c a g g c t g c g t c c a g c a a c a g t g c c c a  
gggctctgatgagtcctc tcatcgct t g t a a (SEQ ID NO:21);

Cw\*030303 :

gctcccac tcca t g a g g t a t t t c t a c a c c g c t g t g t c c c g g c c c g g c c g c g g g a g c c c c a c t t c a t c g c a g t g g g  
c t a c g t g g a c g a c a c g c a g t t c g t g c g g t t c g a c a g c g a c g c c g c g a g t c c g a g a g g g a g c c g c g g g c g c c g t g g  
10 g t g g a g c a g g a g g g c c g g a g t a t t g g g a c c g g g a g a c a c a g a a g t a c a a g c c c a g g c a c a g a c t g a c c g a g t g a  
g c c t g c g g a a c c t g c g c g g c t a c t a c a a c c a g a g c g a g g c c a g g t c t c a c a t c a t c c a g a g g a t g t a t g g c t g c g a  
c g t g g g A c c c g a c g g g c g c c t c c t c c g c g g g t a t g a c c a g t a c g c c t a c g a c g g c a a g g a t t a c a t c g c c c t g a a c  
g a g g a t c t g c g c t c c t g g a c c g c c g g a c a c g g c g g c t c a g a t c a c c c a g c g c a a g t g g g a g g c g g c c c g t g a g g  
c g g a g c a g c t g a g a g c c t a c c t g g a g g g c c t g t g c g t g g a g t g g c t c c g c a g a t a c c t g a a g a a t g g g a a g g a g a c  
15 g c t g c a g c g c g c g g (SEQ ID NO:22);

Cw\*030401 :

a t g c g g g t c a t g g c g c c c c g a a c c c t c a t c c t g c t g c t c t c g g g a g c c c t g g c c c t g a c c g a g a c c t g g g c c g g c t  
c c c a c t c c a t g a g g t a t t t c t a c a c c g c t g t g t c c c g g c c c g g c c g c g g g a g c c c c a c t t c a t c g c a g t g g g c t a  
c g t g g a c g a c a c g c a g t t c g t g c g g t t c g a c a g c g a c g c c g c g a g t c c g a g a g g g a g c c g c g g g c g c c g t g g g t g  
20 g a g c a g g a g g g c c g g a g t a t t g g g a c c g g g a g a c a c a g a a g t a c a a g c c c a g g c a c a g a c t g a c c g a g t g a g c c  
t g c g g a a c c t g c g c g g c t a c t a c a a c c a g a g c g a g g c c g g g t c t c a c a t c A t c c a g a g g a t g t a t g g c t g c g a c g t  
g g g g c c c g a c g g g c g c c t c c t c c g c g g g t a t g a c c a g t a c g c c t a c g a c g g c a a g g a t t a c a t c g c c c t g a a c g a g  
g a t c t g c g c t c c t g g a c c g c c g g a c a c g g c g g c t c a g a t c a c c c a g c g c a a g t g g g a g g c g g c c c g t g a g g c g g  
a g c a g c t g a g a g c c t a c c t g g a g g g c c t g t g c g t g g a g t g g c t c c g c a g a t a c c t g a a g a a t g g g a a g g a g a c g c t  
25 g c a g c g c g c g g a a c a c c c a a g a c a c a c g t g a c c c a c c a t c c c g t c t c t g a c c a t g a g g c c a c c c t g a g g t g c t g g  
g c c c t g g g c t t c t a c c c t g c g g a g a t c a c a c t g a c c t g g c a g t g g g a t g g g g a g g a c c a a a c t c a g g a c a c T g a g c  
t t g t g g a g a c c a g g c c a g c a g g a g a t g g a a c c t t c c a g a a g t g g g c a g c t g t g g t g g t g c c t t c t g g a g a a g a g c a

gagatacacgtgccatgtgcagcacgaggggtgccggagccccacccagagatgggagccgtctccagccc  
accatccccatcgtgggcatcgttgcgtggcctggcgtgtccaggcgtgtccaggagcgtgtgggtggcgtg  
ttgtgatgtltaggaggaagagctcaggtggaaaaggaggagcgtgtctcaggcgtgcgtccagcaacagtgccca  
gggtctgtatgagtcctcctatcgtttgtaa (SEQ ID NO:12);

5 Cw\*030402 :

gtccccactccatgaggtatttctacaccgtgtgtccccggcccgccggggagccccacttcatcgcagtggg  
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gtggagcaggaggggcccggagtattgggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtga  
gccgtcggaacctgcgcggtactacaaccagagcgaggccgggtctcacatcAtccagaggatgtaCggctgcga  
10 cgtggggcccacgggcccctcctccgcggtatgaccagtacgcctacgacggcaaggattacatcgccctgaac  
gaggatctgcgtctcctggaccgcccggacacggcggtcagatcaccagcgcaagtgaggagcgcccggtgagg  
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gctgcagcgcgcg (SEQ ID NO:24);

Cw\*0305 :

15 gtccccactccatgaggtatttctacaccgtgtgtccccggcccgccggggagccccacttcatcgcagtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacccgcgagtcgagaggggagccgcggcgccgtgg  
gtggagcaggaggggcccggagtattgggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtga  
gccgtcggaacctgcgcggtactacaaccagagcgaggccgggtctcacacctccagagCatglacggctgcga  
cgtggggcccacgggcccctcctccgcggtatgaccagtacgcctacgacggcaaggattacatcgccctgaac  
20 gaggatctgcgtctcctggaccgcccggacacggcggtcagatcaccagcgcaagtgaggagcgcccggtgagg  
cggagcagctgagagcctacctggagggcctgtgcgtggagtggtccgcagatacctgaagaaTgggaaggagac  
gctgcagcgcgcg (SEQ ID NO:25);

Cw\*0306 :

25 gtccccactccatgaggtatttctacaccgtgtgtccccggcccgccggggagccccacttcatcgcagtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacccgcgagtcgagaggggagccgcggcgccgtgg  
gtggagcaggaggggcccggagtattgggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtga  
gccgtcggaacctgcgcggtactacaaccagagcgaggccgggtctcacatcctccagaggatglatggctgcga



cgltgggccccgacgggcgccctccctccgcggtatgTccagtiacgcctacgacggcaaggattacatcgccctgaac  
gaggatctgcgtccctggaccgccgacacggcggtcagatcaccagcgcaagtgaggagcgccccgtgagg  
cggagcagctgagagccctacctggagggccctgtgcgtggagtggtccgcagataccigaagaatgggaaggagac  
gctgcagcgcgcg(SAQ ID NO:26);

5 Cw\*0307 :

gtccccactccatgaggtatcttacaccgtgtgtccccggccccggcggggagccccacttcatcgcatggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagccgccccgcccgtg  
gtggagcaggagggcccgagtalgtggaccgggagacacagaagtaagaagccaggcacagactgaccgagtg  
AactgcgaaActgcgcggtacttacaaccagagcgaggccgggtctcacatcAtccagaggatgtatggctgcga  
10 cGiggggccccgacgggcgccctccctccgcggtatgaccagtiacgcctacgacggcaaggattacatcgccctgaac  
gaggatctgcgtccctggaccgccgacacggcggtcagatcaccagcgcaagtgaggagcgccccgtgagg  
cggagcagctgagagccctacctggagggccTgtgcgtggagtggtccgcagataccigaagaatgggaaggagac  
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Cw\*0308 :

15 atgcgggtcalggcccccgaacctcatctgtgtctcgggagccccggccccgaccgagacctgggccccgt  
ccccactccatgaggtatcttacaccgtgtgtccccggccccggcggggagccccacttcatcgcatgggct  
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20 ggggccccgacgggccccctccctccgcggtatgaccagtiacgcctacgacggcaaggattacatcgccctgaacgag  
gatctgcgtccctggaccgccgacacggcggtcagatcaccagcgcaagtgaggagcgccccgtgaggcgg  
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25 tigtggagaccaggccagcaggagatggaacctccagaagtgggcagctgtgtgggtgctcttgagaagagca  
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tigtatgttaggaggaagagctcag (SEQ ID NO:28);

Cw\*0309 :

gctcccatccatgaggtattctacaccgtgtgtccggcccgccgaggagccccacttcacGcagtggg  
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5 gtggagcaggagggccggagtal tgggaccgggagacacagaaglacaaagccaggcacagactgaccgagtga  
gccctgcggaacctgcgcggctactacaaccagagcgaggccgggtctcacatcAtccagaggatgtatggctgcga  
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gaggatctgcgtccctggaccgccgaggacacggcggtcagatcaccagcgcaagtgggagggcgcccgtagg  
cggagcagctgagagcctacctggagggccTgtgcgtggagtggctccgcagatacctgaagaaTgggaaggagac  
10 gctgcagcgcgccg (SEQ ID NO:29);

Cw\*0310 :

gctcccatccatgaggtattctacaccgtgtgtccggcccgccgaggagccccacttcacGcagtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagccgcccgcgcgtgg  
gtggagcaggagggccggagtal tgggaccgggagacacagaaglacaaagccaggcacagactgaccgagtga  
15 gccctgcggaacActgcgcggctactacaaccagagcgaggccgggtctcacatcAtccagaggatgtatggctgcga  
cGtggggcccgacgggcgcctccctccgcggtatgaccagtagccctacgacggcaaggattacatcgccctgaac  
gaggatctgcgtccctggaccgccgaggacacggcggtcagatcaccagcgcaagtgggagggcgcccgtagg  
cggagcagctgagagcctacctggagggccTgtgcgtggagtggctccgcagatacctgaagaaTgggaaggagac  
gctgcagcgcgccg (SEQ ID NO:30);

20 Cw\*0311 :

gctcccatccatgaggtattctacaccgtgtgtccggcccgccgaggagccccgcttcacGcagtggg  
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gtggagcaggagggccggagtal tgggaccgggagacacagaaglacaaagccaggcacagactgaccgagtga  
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25 cgtggggcccgacgggcgcctccctccgcggtatgaccagtagccctacgacggcaaggattacatcgccctgaac  
gaggatctgcgtccctggaccgccgaggacacggcggtcagatcaccagcgcaagtgggagggcgcccgtagg  
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Cw\*0315:

10 gctcccacitccatgaggtatttctacaccgctgtgtcccgcccgccgaggagccccacitcatcgagtgagg  
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15 gaggatctgcgctcttgaccgcccgggacacggcggtcagatcaccagcgcaagtgggaggcgcccggtgagg  
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gctgcagcgcgagg(SEQ ID NO:35);

Cw\*0316:

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20 ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagccgaggcgccgttg  
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25 cggagcagctgagagcctacctggagggcacgtgcgtggagtggctccgcagatacctggagaacgggaaggagac  
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Cw\*040101:

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gagcaggagggggccggagtttgggaccgggagacacagaagtacaagcgccaggcacaggctgaccgagtgaacc  
5 tgcggaaactgcgcggctactacaaccagagcgaggacgggtctcacacctccagaggatgtttggctgcgacct  
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15 gggctctgatgagttctcctatcgtttgtaa (SEQ ID NO:37);  
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25 Cw\*0403 :  
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Cw\*0404 :

15 gctcccactccatgaggatatttctccacatccgtgtccctggcccggccgcggggagccccgcttcatcgagtgagg  
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20 gaggatctgcgctcctggaccgccgaggacacggcggtcagatcaccagcgcaagtgggaggcgcccgtaggg  
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Cw\*0405 :

gctcccactccatgaggatatttctccacatccgtgtccctggcccggccgcggggagccccgcttcatcgagtgagg  
25 ctacCtggacgacacgcagttcgtgcgggttcgacagcgacgccgcgagltccaagaggggagcccggggagccgtgg  
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5 Cw\*0406 :

gctcccaTccatgaggTatttctacaccgctgtgtccggcccgagccgGgagagccccActtcatcgcaTggg  
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Cw\*0407 :

15 gctcccaTccatgaggTatttctccacatccgtgtccTggcccgccgGgggagccccgcttcatcgcaTggg  
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20 gaggatctgcgtccTggaccgccgGgacacggcggtcagatcaccagcgcaagTgggaggcgcccgTgagg  
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Cw\*0408 :

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cc tggggccggacgggcgccttctccgcgggtataaccagttcgcttacgacggcaaggattacatcgccctgaac  
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gctgcagcgcgcgg (SEQ ID NO:44);

5 Cw\*0410 :

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10 cc tggggccggacgggcgccttctccgcgggtataaccagttcgcttacgacggcaaggattacatcgccctgaac  
gaggatctgcgtccctggaccgccgcggacacggcggctcagatcacccagcgcaagtgggaggcgcccgtagg  
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gctgcagcgcgcgg (SEQ ID NO:45);

Cw\*0501 :

15 atgcgggtcatggcggcccgaaacctcatctgtgtctctcgggagccc tggccctgaccgagacctgggccc tgc t  
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gagcaggaggggcccggagtattgggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtgaacc  
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20 ggggcccgcgggcgccttctccgcgggtataaccagttcgcttacgacggcaaggattacatcgccctgaatgag  
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25 ttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtggtgggtgccttctggagaagagca  
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tttgatgttaggaggaagagctcaggtggaaaaggaggagctgcctcaggctgcgtccagcaacagtgccca  
gggctctgaigagtctctcatcgttgttaa (SEQ ID NO:46);

Cw\*0502 :

gtccccacccaatgaggtattttacaccgccgtgtcccgcccgcccgaggagccccgttcatcgcagtggtg  
5 ctacgtggacgacacgcagttcgtgcagttcgacagcgacgcccgagttcaagaggggagcccgggcgccgttg  
gtggagcaggagggcgagttatgggaccgggagacacagaagtacaagcgccaggcacagacigaccgagtga  
acctgcggaaactgcgcggctactacaaccagagcgaggccgggttcacacccctccagaggatgtatggctgcga  
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gaggacctgcgtctctggaccgcccgggacaaggcggtcagatcaccagcgcaagtgggaggcgcccggtgagg  
10 cggagcagcggagagcctaccctggagggtatgtcgtggagtggctGcgagataccctggagaacgggaaggagac  
gttcagcgcgcgg (SEQ ID NO:47);

Cw\*0503 :

atgcgggtcatggcgccccgaacccctcatctgtgtctctcgggagcccggccccgaccgagacctgggctgtct  
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15 cgtggacgacacgcagttcgtgcagttcgacagcgacgcccgagttcaagaggggagcccgggcgccgtgggtg  
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20 agcagcggagagcctaccctggagggtacgtgcgtggagtggctccgcagataccctggagaacgggaagaagacgt  
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25 accatccccatcgtgggcatcgtgtcgtggctggctgtcctggctgtcctagctgtcctaggagctgtgatggctg  
tttgatgttaggaggaagagctcaggtggaaaaggaggagctgcctcaggctgcgtccagcaacagtgccca  
gggctctgaigagtctctcatcgttgttaa (SEQ ID NO:48);

Cw\*0504 :

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gtggagcaggagggccggagtattgggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtga  
5 acctgcggaaactgcgcggctactacaaccagagcgaggccgggtctcacacctccagaggatgtatggctgcga  
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gaggacctgcgtctctggaccgcccgggacaAgcgccgtcagatcacccagcgcaagtgaggagcgcccggtgagg  
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10 Cw\*0505 :

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15 cGtggggcccgacgggcgcctctccgcgggtataaccagtTcgctacgacggcaaggattacatcgccctgaat  
gaggacctgcgtctctggaccgcccgggacaAgcgccgtcagatcacccagcgcaagtgaggagcgcccggtgagg  
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Cw\*0506 :

20 gctccacatccatgaggtatctacacgccgtgtccggcccgccggagagccccgttcatcgcatggg  
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acctgcggaaactgcgcggctactacaaccagagcgaggccgggtctcacacctccagaggatgtatggctgcga  
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25 gaggacctgcgtctctggaccgcccgggacaagcgccgtcagatcacccagcgcaagtgaggagcgcccggtgagg  
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Cw\*0602 :

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5 gaggagggggccggagtattgggaccgggagacacagaagtaagcgccaggcacaggctgaccgagtgaacc  
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15 ttgtgagtgttaggaggaagagctcaggctggaaaaggagggagctgctctcaggctgcgtccagcaacagtgccca  
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Cw\*0603 :

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20 gtggagcaggagggccggagtattgggaccgggagacacagaagtaagcgccaggcacaggctgaccgagtga  
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25 gctgcagcgcgcg(SEQ ID NO:53);

Cw\*0604 :

gctcccacitccatgaggtatctcgacaccgccgtgtcccgcccgcccgaggagagccccgttcatctcagtgagg

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accitcggaatactgcgcggctactacaaccagagcgaggacgggtctcacacccctccagtggaigtatggctgcga  
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5 gaggacctgcgctcctggaccgccgaggacacggcggtcagatcacccagcgcaagtgggagggcgcccgtagg  
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Cw\*0605 :

gtccccactccatgaggtatctcgacaccgccgtgtccggcccgccgaggagccccgttcatctcagtgagg  
10 ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcAagaggggagccCggggcgccgtgg  
gtggagcaggaggggcccggagtatgggaccgggagacacagaagtaacagcgccaggcacaggctgaccgagtga  
accitcggaatactgcgcggctactacaaccagagcgaggacgggtctcacacccctccagtggaigtatggctgcga  
cctggggcccgacgggcgcttctccgcggtatgaccagtcgcctacgacggcaaggattacatcgccctgaac  
gaggacctgcgctcctggaccgccgaggacacGgcgggtcagatcacccagcgcaagtgggagggcgcccgtagg  
15 cggagcagcgagagcctaccitggagggcacgtgcgtggagtggctccgcagataccitggagaacgggaaggagac  
gctgcagcgcgagg (SEQ ID NO:55);

Cw\*0606 :

gtccccactccatgaggtatctcgacaccgccgtgtccggcccgccgaggagccccgttcatctcagtgagg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagccCggggcgccgtgg  
20 gtggagcaggaggggcccggagtatgggaccgggagacacagaagtaacagcgccaggcacaggctgaccgagtga  
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cctggggcccgacgggcgcttctccgcggtatgaccagtcgcctacgacggcaaggattacatcgccctgaac  
gaggacctgcgctcctggaccgccgaggacacggcggtcagatcacccagcgcaagtgggagggcgcccgtagg  
cggagcagcgagagcctaccitggagggcacgtgcgtggagtggctccgcagataccitggagaacgggaaggagac  
25 gctgcagcgcgaggaaacaccaaagacacacgtgaccacacatcccgctcttgaccatgaggccacccitgaggctc  
tggggccctgggtcttaccitcgggagatcacacitgaccitggcagcggtatggcgaggaccaaacitcaggacaccg  
agcttgggagaccaggccagcaggagatggaacctccagaagtgggcagcttgggtggcttcttggagaaga

gcagagat acacgtgccatgtgcagcacgaggggtgccagagccccccacccctgagatgggagccatcttcccag  
cccacatccccatcgtgggcatcgttgcctggccctggctgtccctggctgtccctagctgtccctaggagctgtgAtgg  
ctgttgtgtgtgttaggaggaagagctcag (SEQ ID NO:56);

Cw\*0607 :

5 gctcccactccatgaggtaattcgacaccgccgtgtcccggccccggccgagagccccgcttcatctcagtggg  
ctacgtggacgacacgcagttcgtgcgggtcgacagcgacgcccgagtcgagaggggagccccgggcgccgtgg  
gtggagAaggaggggcccggagtaattgggaccgggagacacagaagtacaagcgccaggcacaggctgaccgagtga  
acctgcggaaactgcgcggctactacaaccagagcgaggacgggtctcacacccctccagtggatgtatggctgcga  
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10 gaggacctgcgtctctggaccgccgagacacggcggtcagatcacccagcgcaagtgggagggcgcccggtgagg  
cggagcagtgagagccctacctggagggcacgtgcgtggagtggctccgcagatacctggagaacgggaaggagac  
gctgcagcgcgagg (SEQ ID NO:57);

Cw\*0608 :

gctcccactccatgaggtaattcgacaccgccgtgtcccggccccggccgagagccccgcttcatctcagtggg  
15 ctacgtggacgacacgcagttcgtgcgggtcgacagcgacgcccgagtcgagaggggagccccgggcgccgtgg  
gtggagcaggaggggcccggagtaattgggaccgggagacacagaagtacaagcgccaggcacaggctgaccgagtga  
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cctggggcccgacgggcgccctccctccgcggtatgaccagtcggcctacgacggcaaggattacatgccccgaac  
gaggacctgcgtctctggaccgccgagacacGgcgggtcagatcacccagcgcaagtgggagggcgcccggtgagg  
20 cggagcagTggagagccctacctggagggcgAgctgcgtggagtggctccgcagatacctggagaacgggaaggagac  
gctgcagcgcgagg (SEQ ID NO:58);

Cw\*0609 :

gctcccactccatgaggtaattcgacaccgccgtgtcccggccccggccgagagccccgcttcatctcagtggg  
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25 gtggagcaggaggggcccggagtaattgggaccgggagacacagaagtacaagcgccaggcacaggctgaccgagtga  
acctgcggaaactgcgcggctactacaaccagagcgaggacgggtctcacacccctccagtggatgtatggctgcga  
cctggggcccgacgggcgccctccctccgcggtataaccagtcggcctacgacggcaaggattacatgccccgaac

gaggacctgcgtccctggaccgccgacacggcggctcagatcaccagcgcaagtgggaggcgcccgtagg  
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gctgcagcgcgcgg(SEQ ID NO:59);

Cw\*070101 :

5 atgcgggtcatggcggcccgagccctccctcctgcgtgcgtcctcgggaggccctggccctgaccgagacctgggctgct  
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cgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagccgcggcgccgtgggtg  
gagcaggagggggccggagtattgggaccgggagacacagaactacaagcgccaggcacaggctgaccgagttagcc  
tgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacacctccagaggatgtAtggctgcgacct  
10 gggggcccgacggcgccctccctccgcggtatgaccagtcgcctacgacggcaaggattacatcgccctgaacgag  
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15 ttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtggtaggtgcttctggacaagagca  
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ctatgatgttaggaggaagagctcaggtaggaaaaggaggagctgtcctcaggctgcgtgcagcaacagtgccea  
gggctctgatgagtcctcctcActtgtaa(SEQ ID NO:60);

20 Cw\*070102 :

atgcgggtcatggcggcccgagccctccctcctgcgtgcgtcctcgggaggccctggccctgaccgagacctgggctgct  
cccacitccatgagglatttcgacaccgccgtgtcccgcccgccgagagccccgcttcatctcagtagggctta  
cgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagccgcggcgccgtgggtg  
gagcaggagggggccggagtattgggaccgggagacacagaactacaagcgccaggcacaggctgaccgagttagcc  
25 tgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacacctccagaggatgtatggctgcgacct  
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agcagctgagagcctaccitggagggcacgtgcgtggagtggtccgcagataccitggagaacgggaaggagacgct  
gcagcgcgcagaacccccaaagacacacgtgaccaccacccccctctctgaccaitaggccacccitagggtgctgg  
gcccitgggtctctacccitgaggagatcacactgacctggcagcgggatggggaggaccagaccaggacaccgagc  
ttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtggtggtgcttctggacaagagca  
5 gagatacacgtgccataitgcagcacgaggggctgcaagagccccctacccitgagctgggagccaatcttcccagccT  
accatccccatcatgggcatcgttgcitggcctggcgttcttgggtgttctttagctgttcttggagctgtggtcaccg  
ctatgatgtgtaggaggaagagctcaggtggaaaaggagggagctgctctcaggctgcgtgcagcaacagtgccca  
gggtctctgatgagtcctctcatcacttgttaa (SEQ ID NO:61);

Cw\*070201 :

10 atgcgggcatggcgccccgagccctcttctgtgtctctcgggaggccitggccctgaccgagacctgggctgtct  
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cgtggacgacacgcagttcgtgcggttgcacagcgacgccgcgagttccgagaggggagccgcggcgccgtgggtg  
gagcaggagggggccggagtattgggaccgggagacacagaagtacaagcgccaggcacaggctgaccgagttagcc  
tgcggaaccttgcgcggctactacaaccagagcgaggacgggtctcacacctccagaggatgttctggctgcgacct  
15 ggggccccgacgggcgccctcttccgcggtatgaccagttccgcttacgacggcaaggattacatcgccccgaacgag  
gacctgcgtctctggaccgcccgggacaccgcggctcagatcacccagcgcaagtTggaggcgccccgtgcggcgg  
agcagctgagagcctaccitggagggcacgtgcgtggagtggtccgcagataccitggagaacgggaaggagacgct  
gcagcgcgcagaacccccaaagacacacgtgaccaccacccccctctctgaccaitaggccacccitagggtgctgg  
gcccitgggtctctacccitgaggagatcacactgacctggcagcgggatggggaggaccagaccaggacaccgagc  
20 ttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtggtggtgcttctggacaagagca  
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ctatgatgtgtaggaggaagagctcaggtggaaaaggagggagctgctctcaggctgcgtgcagcaacagtgccca  
gggtctctgatgagtcctctcatcActtgttaa (SEQ ID NO:62);

25 Cw\*0703 :

tgtctccactccatgaggtatttgcacaccgccgtgtccggccccggcgccggagagccccgttcatctcagitgg  
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ggtaggagcaggagggccggagtaattgggaccgggagacacagaagtacaagcgccaggcacaggcigaccgagtag  
 agccatgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacacctccagaggatgtctggctgcg  
 acctggggcccgacgggcgctccctccgcggtatgaccagtcgcttacgacggcaaggattacatcgccctgaa  
 cgaggacctgcgtctctggaccgcggcgacaccgcggctcagatcacccagcgcaagtagggaggcggcccgtagc  
 5 ggggagcagctagagacctacctggagggactgtgcgtggagtaggctccgcagatacctggagaacgggaaggaga  
 cgtgcagcgcgcaaaccccaaaagacacacgtgaccaccacccccctcttgacctagggccacctgaggtag  
 ctggggccctgggtctctacctgcggagatcacactgacctggcagcgggatggggaggaccagaccaggacacc  
 gagctgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtggtaggtgctcttgacaag  
 agcagagatacacgtgccatatgcagcacgaggggtgcaagagccccctacctgagctgggagccatcttccca  
 10 gccaccatccccatcatgggcatcgtgtctggcctggctgtcttggtgtctctagctgtctctggagctgtggtc  
 accgtatgatgttaggaggaagagctcaggtaggaaaaggaggagctgtctcaggctgcgtgcagcaacagtg  
 ccagggtcttgatgagctctctcatcacttgtaa (SEQ ID NO:63);

Cw\*070401:

atgctgggtcatggcgccccgagccccctctctgtctgtctctcgaggagccctggccctgaccgagacctgggctgtct  
 15 cccacatccatgaggtaattcgacaccgctgtctccggcccgccgcggagagccccgttcalctcagtaggctta  
 cgtggacgacacgcagttcgtgcggttcgacagcgacccgcgagtcggagaggggagccccgggcgctgtgggtg  
 gagcaggaggggcccggagtaattgggaccgggagacacagaagtacaagcgccaggcacaggcigaccgagtagacc  
 tgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacaccttccagaggatgtatggctgcgacct  
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 20 gacctgcgtctctggaccgcccggacaccgcggctcagatcacccagcgcaagtaggagggcccgtagcggcg  
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 25 gagatacacgtgccatatgcagcacgaggggtgcaagagccccctacctgagctgggagccatcttccagccc  
 accatccccatcatgggcatcgtgtctggcctggctgtcttggtgtctctagctgtctctggagctgtggtcaccg  
 ctatgatgttaggaggaagagctcaggtaggaaaaggaggagctgtctcaggctgcgtgcagcaacagtgccca



gggcctctgatgagtcctctcatcActtgtaa (SEQ ID NO:64);

Cw\*070402 :

atgcgggtcatggcgccccgagccctccctccctgcctgcctcgcgggagggcctggccctgaccgagacctgggctgct  
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5 cgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagccccggcgccgtgggtg  
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tgcggaacctgcgcggctactaTaaccagagcgaggacgggtctcacacctccagaggatgtatggctgcgacct  
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gacctgcgctccctggaccgcccgagacaccgcggctcagatcacccagcgcaagttggaggcgcccgctgcggcgg  
10 agcaggacagagcctaccctggaggcgacgtgcgtggagtggctccgcagataccctggagaacgggaagaagacgt  
gcagcgcgcggaacccccaaagacacacgtgacctaccacccccctctctgacctgaggccacctgaggtgctgg  
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tttgggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtgggtggctgcttctggacaagagca  
gagatcacgtgccatacgacgacgaggggtgcaagagccccctacctgagctgggagccatcttccagccc  
15 acctccccatcatgggcatcgttgcctggcctggctgtctgggtgtctctagctgtctctggagctgtggtcacgc  
ctatgatgtgtaggaggaagagctcaggctggaaaaggaggagctgtctctcaggctgcgtgcagcaacagtgccca  
gggcctctgatgagtcctctcatcacttgtaa (SEQ ID NO:65);

Cw\*0705 :

gtccccactccatgaggtatctcgacaccgccgtgtcccgccccggccgagagccccgccttcactcagtggg  
20 ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagcccgggcgccgtgg  
gtggagcaggagggggccgagttatgggaccgggagacacagaagtaagaagccaggcacaggctgaccgagtg  
gccctgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacacctccagaaTaigtatggctgcga  
ccctggggcccgacggcgccctccctcccggggtatgaccagttcgccctacgacggcaaggattacatcgccctgaac  
gaggacctgcgctccctggaccgcccgagacaccgcggctcagatcacccagcgcaagttggaggcgcccgctgcgg  
25 cggagcagctgagagcctaccctggaggcgacgtgcgtggagtggctccgcagataccctggagaacgggaaggagac  
gtgcagcgcgag (SEQ ID NO:66);

Cw\*0706 :

a t g c g g g t c a t g g c g c c c g a g c c c t c c t c c t g c t g c t c t c g g g a g g c c t g g c c c t g a c c g a g a c c t g g g c c t g c t  
c c c a c t c c a t g a g g t a t t t c g a c a c c g c c g t g t c c c g g c c c g g c c g g a g a g c c c c g c t t c a t c t c a g t g g g c t a  
c g t g g a c g a c a c g c a g t t c g t g c g g t t c g a c a g c g a c g c c g c g a g t c c g a g a g g g g a g c c g c g g g c c c g t g g g t g  
g a g c a g g a g g g g c c g g a g t a t t g g g a c c g g g a g a c a c a g a a c t a c a a g c g c c a g g c a c a g g c t g a c c g a g t g a g c c  
5 t g c g g a a c c t g c g c g g c t a c t a c a a c c a g a g c g a g g a c g g g t c t c a c a c c c t c c a g a g g a t g t a t g g c t g c g a c c t  
g g g g c c c g a c g g g c g c c t c c t c c g c g g g t a t g a c c a g t c c g c c t a c g a c g g c a a g g a t t a c a t c g c c c t g a a c g a g  
g a c c t g c g c t c c t g g a c c g c c g g a c a c c g c g g c t c a g a t c a c c c a g c g c a a g t t g g a g g c g g c c c g t g c g g c g g  
a g c a g c t g a g a g c c t a c c t g g a g g g c a c g t g c g t g g a g t g g c t c c g c a g a t a c c t g g a g a a c g g g a a g g a g a c g c t  
g c a g c g c g c a g a a c c c c a a a g a c a c a c g t g a c c c a c c a c c c c t c t c t g a c c a t g a g g c c a c c c t g a g g t g c t g g  
10 g c c c t g g g c t t c t a c c c t g c g g a g a t c a c a c t g a c c t g g c a g c g g g a t g g g g a g g a c c a g a c c c a g g a c a c c g a g c  
t t g t g g a g a c c a g g c c a g c a g g a g a t g g a a c c t t c c a g a a g t g g g c a g c t g t g g t g g t g c c t t c t g g a c a a g a g c a  
g a g a t a c a c g t g c c a t a t g c a g c a c g a g g g g c t g c a a g a g c c c c t c a c c c t g a g c t g g g a g c c a t c t t c c c a g c c c  
a c c a t c c c c a t c a t g g g c a t c g t t g c t g g c c t g g c t g t c c t g g t t g t c c t a g c t g t c c t t g g a g c t g t g g t c a c c g  
c t a A g a t g t g t a g g a g a a g a g c t c a g g t g g a a a a g g a g g g a g c t g c t c t c a g g t t g c g t g c a g c a a c a g t g c c c a  
15 g g g c t c t g a t g a g t c t c t c a t c a c t t g l a a (SEQ ID NO:67);

Cw\*0707 :

g c t c c c a c t c c a t g a g g t a t t t c g a c a c c g c c g t g t c c c g g c c c g g c c g g a g a g c c c c g c t t c a t c t c a g t g g g  
c t a c g t g g a c g a c a c g c a g t t c g t g c g g t t c g a c a g c g a c g c c g c g a g t c c g a g a g g g a g c c g c g g g c c c g t g g  
g t g g a g c a g g a g g g g c c g g a g t a t t g g g a c c g g g a g a c a c a g a a c t a c a a g c g c c a g g c a c a g g c t g a c c g a g t g a  
20 a c c t g c g g a a A c t g c g c g g c t a c t a c a a c c a g a g c g a g g c c g g g t c t c a c a c c c t c c a g a g g a t g t A t g g c t g c g a  
c c t g g g g c c c g a c g g g c g c c t c c t c c g c g g g t a t g a c c a g t C g c c t a c g a c g g c a a g g a t t a c a t c g c c c t g a a c  
g a g g a c c t g c g c t c c t g g a c c g c c g g a c a c c g c g g c t c a g a t c a c c c a g c g c a a g t T g g a g g c g g c c c g t g c g g  
c g g a g c a g c t g a g a g c c t a c c t g g a g g g c a c g t g c g t g g a g t g g c t c c g c a g a t a c c t g g a g a a c g g g a a g g a g a c  
g c t g c a g c g c g A g (SEQ ID NO:68);

25 Cw\*0708 :

g c t c c c a c t c c a t g a g g t a t t t c g a c a c c g c c g t g t c c c g g c c c g g c c g g a g a g c c c c g c t t c a t c t c a g t g g g  
c t a c g t g g a c g a c a c g c a g t t c g t g c g g t t c g a c a g c g a c g c c g c g a g t c c g a g a g g g a g c c g c g g g c c c g t g g

gtggagcaggaggggcccggagtattgggaccgggagacacagaagtacaagcgccaggcacaggctgaccgagtga  
gacctgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacacctccagaggatgtTggctgcga  
cctggggcccgacgggcgctcctccgcggtatgaccagTCgcctacgacggcaaggattacatcgccctgaac  
gaggacctgcgctcctggaccgcccggacaccgcggtcagatcaccagcgcaagtTggaggcggcccgctgcgg  
5 cggagcagctgagagcctacctggagggcacgtgcgtggagtggctccgcagataccTggagaacgggaaggagac  
gctgcagcgcgAg (SEQ ID NO:69);

Cw\*0709 :

gtccctccatgaggtaattcgacaccgcccgtgtccggcccggccgaggagccccgttcaatctcagtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagccgcccgcgcgtgg  
10 gtggagcaggaggggcccggagtattgggaccgggagacacagaactacaagcgccaggcacaggctgaccgagtga  
acctgcggaAActgcgcggctactacaaccagagcgaggAcgggtctcacacctccagaggatgtAtggctgcga  
cctggggcccgacgggcgctcctccgcggtatgaccagTCgcctacgacggcaaggattacatcgccctgaac  
gaggacctgcgctcctggaccgcccggacaccgcggtcagatcaccagcgcaagtTggaggcggcccgctgcgg  
cggagcagctgagagcctacctggagggcacgtgcgtggagtggctccgcagataccTggagaacgggaaggagac  
15 gctgcagcgcgAg (SEQ ID NO:70);

Cw\*0710 :

gtccctccatgaggtaattcgacaccgcccgtgtccggcccggccgaggagccccgttcaatctcagtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagccgcccgcgcgtgg  
gtggagcaggaggggcccggagtattgggaccgggagacacagaagtacaagcgccaggcacaggctgaccgagtga  
20 gacctgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacatcAtccagaggatgtCtggctgcga  
cctggggcccgacgggcgctcctccgcggtatgaccagTCgcctacgacggcaaggattacatcgccctgaac  
gaggacctgcgctcctggaccgcccggacaccgcggtcagatcaccagcgcaagtTggaggcggcccgctgcgg  
cggagcagctgagagcctacctggagggcacgtgcgtggagtggctccgcagataccTggagaacgggaaggagac  
gctgcagcgcgAg (SEQ ID NO:71);

25 Cw\*0711 :

atgcgggtcatggcgccccgagccctcctcctgtcgtcctcgggaggcctggccctgaccgagacctgggccctgt  
ccctccatgaggtaattcgacaccgcccgtgtccggcccggccgaggagccccgttcaatctcagtgggcta

cgtaggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagccccgggcgccgtgggtg  
gagcaggagggggccggagtaattgggaccgggagacacagaagtaacaagcgccaggcacaggctgaccgagtagacc  
tgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacacctccagaggatgtatggctgcgacct  
ggggcccgacgggcgctctctccgcggtatgaccagttcgctacgacggcaaggattacatcgccctgaacgag  
5 gacctgcgtctctggaccgcccgggacaccgcggctcagatcaccagcgcaagttggaggcgcccgtagcgcg  
agcaggaCagagcctacctggagggcacgtgcgtggagtaggctccgcagatactggagaacgggaagaagacgt  
gcagcgcgcggaacccccaaagacacacgtgaccaccacccccctctctgacctaggccaccttaggtgctgg  
gcccagggtctctacctgcggagatcacactgacctggcagcggtatggggaggaccagaccaggacaccgagc  
ttgtggagaccaggccagcaggagatggaacctccagaagtgggcagctgtggtaggtgctctctggacaagagca  
10 gagatacacgtgccataatgcagcacgaggggctgcaagagccccctacctgagctgggagccatctctccagccc  
accatccccatcatgggcatcgttgcgtggcctggctgtcttggtgtctctagctgtctctggagctgtggtcaccg  
ctatgatgttaggaggaagagctcaggtggaaaaggaggagctgctctcaggctgcgtGcagcaacagtgccca  
gggtctgtatgagctctctcatcgcttgtaa (SEQ ID NO:72);

Cw\*0712 :

15 gctcccatccatgaggtatttcgacaccgccgtgtccggccccggcgaggagccccgcttcatctcagtggg  
ctactggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagccccgggcgccgtgg  
gtggagcaggagggggccggagtaattgggaccgggagacacagaagtaacaagcgccaggcacaggctgaccgagtaga  
gccitgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacacctccagaggatgtatggctgcga  
cctggggcccgacgggcgctctctccgcggtatgaccagttcgctacgacggcaaggattacatcgccctgaac  
20 gaggacctgcgtctctggaccgcccgggacaccgcggctcagatcaccagcgcaagttggaggcgcccgtagcgg  
cggagcaggaCagagcctacctggagggcacgtgcgtggagtaggctccgcagatactggagaacgggaagaagac  
gctgcagcgcgcg (SEQ ID NO:73);

Cw\*0713 :

gctcccatccatgaggtatttcgacaccgccgtgtccggccccggcgaggagccccgcttcatctcagtggg  
25 ctactggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagcccggggcgccgtgg  
gtggagcaggagggggccggagtaattgggaccgggagacacagaagtaacaagcgccaggcacaggctgaccgagtaga  
gccitgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacacctccagaggatgtctggctgcga

ccctggggcccgacgggcgcctccctccgcgggatgaccagTcgccctacgacggcaaggattacatcgccctgaac  
gaggacctgcgctccctggaccgccgcggacaccgcggctcagatcacccagcgcaagtTggaggcggcccgctgcgg  
cggagcagctgagagcctaccctggagggcacgtgcgtggagtggtccgcagataccctggagaacgggaaggagac  
gctgcagcgcgAg (SEQ ID NO:74);

5 Cw\*0714 :

gctcccactccatgaggtatttcgacaccgccgtgtcccgcccgccgcggagagccccgcttcaatcagtgagg  
ctacgtggacgacacgcagttcgctgcggttcgacagcgacgccgcgagtcgagaggggagccgcgggcgcctgg  
gtggagcaggaggggcccggagatctgggaccgggagacacagaagtaagcggccaggcacaggctgaccgagtga  
gccctgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacacctccagaggatgtacggctgcga

10 ccctggggcccgacgggcgcctccctccgcgggatgaccagtcgcctacgacggcaaggattacatcgccctgaac  
gaggacctgcgctccctggaccgccgcggacaccgcggctcagatcacccagcgcaagtTggaggcggcccgctgcgg  
cggagcagctgagagcctaccctggagggcacgtgcgtggagtggtccgcagataccctggagaacgggaaggagac  
gctgcagcgAcag (SEQ ID NO:75);

Cw\*0715 :

15 gctcccactccatgaggtatttcgacaccgccgtgtcccgcccgccgcggagagccccgcttcaatcagtgagg  
ctacgtggacgacacgcagttcgctgcggttcgacagcgacgccgcgagtcgagaggggagccgcgggcgcctgg  
gtggagcaggaggggcccggagatctgggaccgggagacacagaagtaagcggccaggcacaggctgaccgagtga  
gccctgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacacctccagaggatgtctggctgcga  
ccctggggcccgacgggcgcctccctccgcgggatgaccagtcgcctacgacggcaaggattacatcgccctgaac

20 gaggacctgcgctccctggaccgccgcggacaccgcggctcagatcacccagcgcaagtTggaggcggcccgctgcgg  
cggagcagctgagagcctaccctggagggcAgctgcgtggagtggtccgcagataccctggagaacgggaaggagac  
gctgcagcgcgAg (SEQ ID NO:76);

Cw\*0716 :

gctcccactccatgaggtatttcgacaccgccgtgtcccgcccgccgcggagagccccgcttcaatcagtgagg

25 ctacgtggacgacacgcagttcgctgcggttcgacagcgacgccgcgagtcgagaggggagccgcgggcgcctgg  
gtggagcaggaggggcccggagatctgggaccgggagacacagaatcaagcggccaggcacaggctgaccgagtga  
gccctgcggaacctgcgcggctactacaaccagagcgaggcgggtctcacacctccagaggatgtatggctgcga

ccctggggcccgacgggcgcctccctccgcggtatgaccagtcgcctacgacggcaaggattacatcgccctgaac  
gaggacctgcgcctcctggaccgcccgggacaccgcggctcagatcaccagcgcaagttggaggcgcccgctgcgg  
cggagcagctgagagcctaccctggagggcacgtgcgtggagtggctccgcagataccctggagaacgggaaggagac  
gctgcagcgcgcAg (SEQ ID NO:77);

5 Cw#0717:

gctcccactccatgaggtatttcgacaccgccgtgtcccgcccgcccgaggagccccgcctcaictcagtggg  
ctacgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagtcgagaggggagcccgggcgccgtgg  
gtggagcaggagggcgaggatattgggaccgggagacacagaagtaagcggccaggcacaggctgaccgagtga  
gcctgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacacctccagaggatgtctggctgcga

10 cctggggcccgacgggcgcctccctccgcggtatgaccagtcgcctacgacggcaaggattacatcgccctgaac  
gaggacctgcgcctcctggaccgcccgggacaccgcggctcagatcaccagcgcaagttggaggcgcccgctgcgg  
cggagcagctgagagcctaccctggagggcacgtgcgtggagtggctccgcagataccctggagaacgggaaggagac  
gctgcagcgcgcagaacccccaaagacacacgtgaccaccacccccctctctgaccatgaggccacctgaggtgc  
tgggcccctgggcttctacccctgcggagatcacactgacctggcagcgggatggggaggaccagaccaggacaccg  
15 agcttgtggagaccaggccagcaggagatggaacctccagaagtgggcagctgttggtggcttcttgacaaga  
gcagagatacacgtgccatagtcagcacgaggggctgcaagagccccctacctgagctggg (SEQ ID

NO:78);

Cw#0718:

atgcgggtcaltggcgccccgagccctccctcctgtctgtctcgggaggccctggccctgaccgagacctgggctgct  
20 cccacitccatgaggtatttcgacaccgccgtgtcccgcccgcccgaggagccccgccttcatctcagtgggcta  
cgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagtcgagaggggagcccgggcgccgtgggtg  
gagcaggagggcgaggatattgggaccgggagacacagaactacaagcggccaggcacaggctgaccgagttagcc  
tgcggaacctgcgcggctactacaaccagagcgaggacgggtctcacacctccagaggatgtatggctgcgacct  
ggggcccgacgggcgcctccctccgcggtatgaccagtcgcctacgacggcaaggattacatcgccctgaacgag  
25 gacctgcgcctcctggaccgcccgggacaccgcggctcagatcaccagcgcaagttggaggcgcccgctgcggcgg  
agcagctgagagcctaccctggagggcacgtgcgtggagtggctccgcagataccctggagaacgggaaggagacgt  
gcagcgcgcagaacccccaaagacacacgtgaccaccacccccctctctgaccatgaggccacctgaggtgctgg

gccc tgggt t t t t accc t g c g g a g a t c a c a c t g a c c t g g c a g c g g g a t g g g g a g g a c c a g a c c c a g g a c a c c g a g c  
t t g t g g a g a c c a g g c c a g c a g g a g a t g g a a c c t t c c a g a a g t g g g c a g c t g t g g t g g t g c c t t c t g g a c a a g a g c a  
g a g a t a c a c g t g c c a t a t g c a g c a c g a g g g c t g c a a g a c c c c t a c c c t g a g c t g g g a g c c a t c t t c c a g c c c  
a c c a t c c c c a t c a t g g g c a t c g t t g c t g g c c t g g c t g t c c t g g t t g t c c t a g c t g t c c t t g g a g c t g t g g t c a c c g  
5 c t a t g a t g t g t a g g a g a a g a g c t c a g g t g g a a a g g a g g g a g c t g c t c t c a g g t g c g t g c a g c a a c a g t g c c c a  
g g g c t c t g a t g a g t c t c t c a t c a c t t g t a a (SEQ ID NO:79);

Cw\*080101 :

a t g c g g g t c a t g g c g c c c g a a c c c t c a t c c t g c t g c t c t c g g g a g c c c t g g c c c t g a c c g a g a c c t g g g c c t g c t  
c c c a c t c c a t g a g g t a t t t c t a c a c c g c c g t g t c c c g g c c g g c c g g a g a g c c c c g t t c a t c g c a g t g g g c t a  
10 c g t g g a c g a c a c g c a g t t c g t g c a g t t c g a c a g c g a c g c c g c g a g t c c a a g a g g g g a g c c g c g g g c g c c g t g g g t g  
g a g c a g g a g g g g c c g g a g t a t t g g g a c c g g g a g a c a c a g a a g t a c a a g c g c c a g g c a c a g a c t g a c c g a g t g a g c c  
t g c g g a a c c t g c g c g g c t a c t a c a a c c a g a g c g a g g c c g g g t c t c a c a c c c t c c a g a g g a t g t a t g g c t g c g a c c t  
g g g g c c c g a c g g g c g c c t c c t c c g c g g g t a t a a c c a g t t c g c c t a c g a c g g c a a g g a t t a c a t c g c c c t g a a t g a g  
g a c c t g c g c t c c t g g a c c g c c g c g g a c a c g g c g g c t c a g a t c a c c c a g c g c a a g t g g g a g g c g g c c c g t a c g g c g g  
15 a g c a g c t g a g a g c c t a c c t g g a g g g c a c g t g c g t g g a g t g g c t c c g c a g a t a c c t g g a g a a c g g g a a g a a g a c g c t  
g c a g c g c g c g g a a c a c c c a a g a c a c a c g t g a c c c a c c a t c c c g t c t c t g a c c a t g a g g c c a c c c t g a g g t g c t g g  
g c c c t g g g c t t c t a c c c t g c g g a g a t c a c a c t g a c c t g g c a g c g g g a t g g c g a g g a c c a a a c t a g g a c a c c g a g c  
t t g t g g a g a c c a g g c c a g c a g g a g a t g g a a c c t t c c a g a a g t g g g c a g c t g t g g t g g t g c c t t c t g g a g a a g a g c a  
g a g a t a c a c g t g c c a t g t g c a g c a c g a g g g c t g c c a g a g c c c c t a c c c t g a g a t g g g G g c c a t c t t c c a g c c c  
20 a c c a t c c c c a t c g t g g g c a t c g t t g c t g g c c t g g c t g t c c t g g c t g t c c t a g c t g t c c t a g g a g c t g t g a t g g c t g  
t t g t g a t g t g t a g g a g a a g a g c t c a g g t g g a a a g g a g g g a g c t g c t c t c a g g c t g c g t c c a g c a a c a g t g c c c a  
g g g c t c t g a t g a g t c t c t c a t c g c t t g t a a (SEQ ID NO:80);

Cw\*080102 :

g c t c c c a c t c c a t g a g g t a t t t c t a c a c c g c c g t g t c c c g g c c g g c c g g a g a g c c c c g t t c a t c g c a g t g g g  
25 c t a c g t g g a c g a c a c g c a g t t c g t g c a g t t c g a c a g c g a c g c c g c g a g t c c a a g a g g g g a g c c g c g g g c g c c g t g g  
g t g g a g c a g g a g g g g c c g g a g t a t t g g g a c c g g g a g a c a c a g a a g t a c a a g c g c c a g g c a c a g a c t g a c c g a g t g a  
g c c t g c g g a a c c t g c g c g g c t a c t a c a a c c a g a g c g a g g c c g g g t c t c a c a c c c t c c a g a g g a t g t a C g g c t g c g a

cttggggcccgacgggcgccttctccgcgggtataaaccagtTcgcttacgacggcaaggattacatcgccctgaat  
gaggacctgcgttcttgaccgccgcggacacggcggctcagatcaccagcgcaagtgggaggcgcccgTAcgg  
cggagcagctgagagcctacctggagggcacgtgcgtggagtggtccgcagatacctggagaacgggaagaagac  
gctgcagcgcgcg(SAQ ID NO:81);

## 5 Cw\*0802 :

atgcgggtcatggcggccgaacccatcttgcgtctcgggagccctggccctgaccgagacctgggcttgc  
cccactccaatgaggtatttctacaccgccgtgtccggcccgccggagagccccgttcatcgagtggtta  
cgtggacgacacgcagttcgtgcagttcgacagcgacgccgcgagtcgaagaggggagccgcgggcgcctgggtg  
gagcaggaggggcccggagttatgggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtgagcc  
10 tgcggaacctgcgcggctactacaaccagagcgaggccgggtctcacacctccagaggatglatggctgcgacct  
ggggcccgacgggcgccttctccgcgggtataaaccagtTcgcttacgacggcaaggattacatcgccctgaatgag  
gacctgcgttcttgaccgccgcggacaAggcggctcagatcaccagcgcaagtgggaggcgcccgtagggcgg  
agcagcggagagcctacctggagggcacgtgcgtggagtggtccgcagatacctggagaacgggaagaagacgct  
gcagcgcgcggaacacccaaagacacacgtgaccacatcccgtctctgacctgaggccacctgagggtgc  
15 gcccgggcttctacctgcggagatcacactgacctggcagcgggatggcgaggacaaactcaggacaccgagc  
ttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtgggtggcttcttggagaagagca  
gagatcacgtgccatgtgcagcacgaggggtgccagagccccacacctgagatgggGgccaatcttccagccc  
accatccccatcgtgggcatcgttgcctggctggctgtcttggctgtcttagctgtcttaggagctgtgatggctg  
ttgtgatgtgtaggaggaagagctcagggtgaaaaggaggagctgtctcaggctgcgtccagcaacagtgccca  
20 gggctctgatgagtcctcatcgcttggtaa(SAQ ID NO:82);

## Cw\*0803 :

atgcgggtcatggcggccgaacccatcttgcgtctcgggagccctggccctgaccgagacctgggcttgc  
cccactccaatgaggtatttctacaccgccgtgtccggcccgccggagagccccgttcatcgagtggtta  
cgtggacgacacgcagttcgtgcagttcgacagcgacgccgcgagtcgaagaggggagccgcgggcgcctgggtg  
25 gagcaggaggggcccggagttatgggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtgagcc  
tgcggaacctgcgcggctactacaaccagagcgaggccgggtctcacacctccagaggatglatggctgcgacct  
ggggcccgacgggcgccttctccgcgggtataaaccagtTcgcttacgacggcaaggattacatcgccctgaatgag



gacctgcgctccctggaccgccgacacggcggctcagatcaccagcgcaagtgggaggcgcccgctacggcgg  
agcagctgagagcctaccctggagggcacgtgcgtggagtggctccgcagataccctggagaacAggaagaagacgct  
gcagcgcgcggaacacccaaagacacacgtgaccaccatcccgtctctgacatgaggccacccctgaggctgctgg  
gcccctgggcttctacccctgcggagatcacactgacctggcagcgggatggcgaggaccaaactcaggacaccgagc  
5 tttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtggctggcttctggagaagagca  
gagatcacgctgccaatgtgcagcacgaggggctgccagagccccctacccctgagatgggggccaatcttccagccc  
accatccccatcgtgggcatcgttgcctggcctggcgttccctaggctgtcctaggagctgtgAtggcgtg  
ttgtgagtgttaggaggaagagctcaggctggaaaaggaggaggcgtgcctcaggctgcgtccagcaacagtgccca  
gggctctgagtgcgtctctcatcgcttgtaa (SEQ ID NO:83);

10 Cw\*0804 :

gctccactccatgaggtatttctacaccgccgtgtcccgcccgccgagagccccgcttcatcgcagtggg  
ctacgtggacgacacgcagttcgtgcagttcgacagcgacgcccgagttcaagaggggagcccgggcgccgtgg  
gtggagcaggaggggcccggagtattgggaccgggagacacagaagtlacaagcgccaggcacagactgaccgagtga  
gacctgcggaacctgcgcggctactacaaccagagcgaggccgggtctcacacccctccagaggatgtatggctgcga  
15 cctggggcccgacggcgccctctcccggggtataaccagttcgccctacgacggcaaggattacatgccccgaat  
gaggacctgcgctccctggaccgccgaggacaAggcggctcagatcaccagcgcaagtgggaggcgcccgctgagg  
cggagcagcTgagagcctaccctggagggcacgtgcgtggagtggctccgcagataccctggagaacgggaagAagac  
gctgcagcgcgagg (SEQ ID NO:84);

Cw\*0805 :

20 gctccactccatgaggtatttctacaccgccgtgtcccgcccgccgagagccccgcttcatcgcagtggg  
ctacgtggacgacacgcagttcgtgcagttcgacagcgacgcccgagttcaagaggggagcccgggcgccgtgg  
gtggagcaggaggggcccggagtattgggaccgggagacacagaagtlacaagcgccaggcacagGctgaccgagtga  
gacctgcggaacctgcgcggctactacaaccagagcgaggccgggtctcacacccctccagaggatgtatggctgcga  
cctggggcccgacggcgccctctcccggggtataaccagttcgccctacgacggcaaggattacatgccccgaat  
25 gaggacctgcgctccctggaccgccgaggacaAggcggctcagatcaccagcgcaagtgggaggcgcccgctgagg  
cggagcagcgagagcctaccctggagggcacgtgcgtggagtggctccgcagataccctggagaacgggaagAagac  
gctgcagcgcgagg (SEQ ID NO:85);

Cw\*0806 :

gctcccactccatgaggtatttctacaccgccgtgtcccgcccgccgagagccccgccttcalcgagtgagg  
ctacgtggacgacacgcagttcgtgcagttcgacagcgacgccgcgagttccaagaggggagccgcggcgccgtgg  
gtggagcaggagggggccggagtattgggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtga  
5 gacctgcggaacctgcgcggctactacaaccagagcgaggccgggtctcacacccctccagaggatgtatggctgcga  
cctggggcccgacgggcgcttctccgcggtataaccagttcgcttacgacggcaaggattacatcgccctgaat  
gaggacctgcgcttctggaccgccgagacacggcggtcagatcacccagcgcaagtgggaggcgcccgctacgg  
cggagcagctgagagcctacctggagggcGcgtgcgtggagtggctccgcagatacctggagaacAggaagaagac  
gctgcagcgcgcg (SEQ ID NO:86);

10 Cw\*0807 :

gctcccactccatgaggtatttctacaccgccgtgtcccgcccgccgagagccccgccttcalcgagtgagg  
ctacgtggacgacacgcagttcgtgcagttcgacagcgacgccgcgagttccaagaggggagccgcggcgccgtgg  
gtggagcaggagggggccggagtattgggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtga  
gacctgcggaacctgcgcggctactacaaccagagcgaggccgggtctcacacccctccagaggatgtatggctgcga  
15 cctggggcccgacgggcgcttctccgcggtataaccagttcgcttacgacggcaaggattacatcgccctgaat  
gaggacctgcgcttctggaccgccgagacaAggcggctcagatcacccagcgcaagtTggaggcgcccgctgagg  
cggagcagcgagagcctacctggagggcacgtgcgtggagtggctccgcagatacctggagaacgggaagAagac  
gctgcagcgcgcg (SEQ ID NO:87);

Cw\*0808 :

20 gctcccactccatgaggtatttctacaccgccgtgtcccgcccgccgagagccccgccttcalcgagtgagg  
ctacgtggacgacacgcagttcgtgcagttcgacagcgacgccgcgagttccaagaggggagccgcggcgccgtgg  
gtggagcaggagggggccggagtattgggaccgggagacacagaagtacaagcgccaggcacagactgaccgagtga  
gacctgcggaacctgcgcggctactacaaccagagcgaggccgggtctcacacccctccagagCatgtatggctgcga  
cctggggcccgacgggcgcttctccgcggtataaccagttcgcttacgacggcaaggattacatcgccctgaat  
25 gaggacctgcgcttctggaccgccgagacacggcggtcagatcacccagcgcaagtgggaggcgcccgctacgg  
cggagcagctgagagcctacctggagggcacgtgcgtggagtggctccgcagatacctggagaacgggaagAagac  
gctgcagcgcgcg (SEQ ID NO:88);

Cw\*0809 :

atgcgggtcatggcgccccgaacccctacccctgctgctcgcggagcccggcccagaccagaccctgggctgct  
cccactccaatgaggtatctctacaccgcccgtgtcccgccccggccgagagccccgcttcatcgcagtgggctta  
cgtggacgacacgcagttcgtgcagttcgacagcgacgccgcgagttcaagaggggagccgcccggcgccgtgggtg  
5 gaggcaggaggggcccggagtattgggaccgggagacacagaagtaagaagccaggcacagactgaccgagttagcc  
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10 gcagcgcgcggaacacccaaagacacacgtgaccacccatcccgtctctgacctgaggccacctgaggtgctgg  
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Cw\*120201 :

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15 cgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagttcaagaggggagccgcccggcgccgtgggtg  
gaggcaggaggggcccggagtattgggaccgggagacacagaagtaagaagccaggcacaggtgaccgagttagcc  
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25 accatccccatcgtgggcatcgtgtcgtggctggctgtcctggctgtcctagctgtcctaggagctgtgAtggctg  
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Cw\*120202 :

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cgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagttccaagaggggagccgccccgcccgtgggtg  
5 gagcaggaggggcccggaglatgggaccgggagacacagaagtlacaagcggcagggcacaggctgaccgagttagcc  
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10 gcagcgcgcggaacacccaaagacacacgtgaccacccatcccgtctctgacctgagggccacctgaggtgctgg  
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15 ttgtgatgtgtaggaggaagagctcaggtaggaaaaggagggagctgctctcaggctgcgtccagcaacagtgccca  
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Cw\*120203 :

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25 gctgcagcgcgcg (SEQ ID NO:92);

Cw\*120301 :

atgcgggtcatggcgccccgaacctcatcctgctgctctcgggagccctggccctgaccgagacctgggctgct

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5 ggggccccgacggcgccctctccgcgggtatgaccagtcgccctacgacggcaaggattacatcgccccgaacgag  
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10 ttgtggagaccagccagcaggagatggaaccttccagaagtgggcagctgtgggggtgcttctggagaagagca  
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15 Cw\*120302 :

gctcccac tccatgaggtatttctacaccgccgtgtccggcccgccggagagccccgttca tgcagtggg  
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gtggagcaggagggggccggagttatgggaccgggagacacagaagta caagcgccaggcacaggctgaccgagtga  
gctgcggaacctgcgcggctactacaaccagagcgaggccgggtctcacacctccagtggatglatggctgcga  
20 cctggggcccgacggcgccctctccgcgggtatgaccagtcgccctacgacggcaaggattacatcgccccgaac  
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Cw\*120401 :

25 atgggggtcatggcgccccgaaccttca tccgtgtctctcgggagccccggccccgaccgagacctgggctgt  
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442

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5 agcagTggagagcctacctggagggcacgtgcgtggagtggtccgcagatacctggagaacgggaaggagacgt  
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Q ID NO:95);

Cw\*120402 :

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10 cccactccatgaggtattctacaccgcccgtctccggcccggccgagagccccgcttcatcgcatgggcta  
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20 gagatacacgtgccatgtgcagcacgaggggtgccagagccccacccctgagatgggagccatcttccagccc  
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ttgtgaltgttaggaggaagagctcaggtggaaaaggaggagctgtctctcaggtgcgtccagcaacagtgccca  
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Cw\*1205 :

25 atgcggtcatggcgccccgaacctcatctctgtctctcgaggagccctggccctgaccgagacctgggctgct  
cccactccatgaggtattctacaccgcccgtctccggcccggccgagagccccgcttcatcgcatgggcta  
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5 agcagtgagagccctaccctggaggcacgtgcgtggagtggctccgcagataccctggagaacgggaaggagacgt  
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10 accatccccatcgctgggcatcgctgcctggctggctgtcttggctgtctttagctgtctttaggagctgtgAtggctg  
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Cw\*1206 :

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15 ctacttgagacacacgcagttcgtgcggctcgacagcgacgccgcgagttcaagaggggagccgcgggcgcccgtgg  
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20 cggagcagtgagagccctaccctggagggcacgtgcgtggagtggctccgcagataccctggagaacgggaaggagac  
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Cw\*1207 :

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25 gtggagcaggaggggcccggagtal tgggaccgggagacacagaaglacaaagcggcaggcacaggctgaccgagtG  
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Cw\*1208:

5 atgcgggicattggcggccgaacccatccctgctgctctcgggagccctggccctgaccgagacctgggctgct  
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10 ggggcccagggcgccctccctccgcggtatgaccagtcgcccacgacggcaaggattacatcgccctgaacgag  
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15 ttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtgtgtgtgcttctggagaagagca  
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20 Cw\*140201:

atgcgggicattggcggccgaacccatccctgctgctctcgggagccctggccctgaccgagacctgggctgct  
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cgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagttccgagaggggagccgcggcgccgtgggtg  
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25 tgcggaacctgcgcggctactacaaccagagcgaggccgggtctcacacctccagtaggttggctgcgacct  
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5 gagatacacgtgccatgtgcagcacgaggggtgccggagcccc tacc tggatgggagccgtcttcccagccc  
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ttgtgatgtgtaggaggaagagctcagg tggaaaaggagggagctgtctcaggctgcgtccagcaacagtgccca  
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Cw\*140202 :

10 gctcccactccatgaggtatttctCcacatccgtgtcccggccccggccgaggagccccgttcatcgcagtaggg  
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gtggagcaggaggggcccggaglat tgggaccgggagacacagaagtacaagcgccaggcacagac taccgagtgga  
gcc tgcggaacctgcgcggctactacaaccagagcgaggccgggtctcacacctccag tggatgtTtggctgcga  
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15 gaggacctgcgtctctggaccgcccgggacacggcggctcagatcacccagcgcaagtgggagggcgcccggtgagg  
cggagcagcggagagcc tacc tggagggcacgtgcgtggagtaggc tccgcagat acc tggagaacgggaaggagac  
gctgcagcgcgcgg (SEQ ID NO:102);

Cw\*1403 :

atgcgggicattggcgccccgaacctcatctctgtctctcgggagcccc tggccc taccgagaccttggccctgct  
20 cccactccatgaggtatttctCcacatccgtgtcccggccccggccgaggagccccacttcatcgcagtagggcta  
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gccc tgggc t t c t a c c c t g c g g a g a t c a c a c t g a c c t g g c a g t g g g a t g g g g a g g a c c a a a c t c a g g a c a c c g a g c  
t t g t g g a g a c c a g g c c a g c a g g a g a t g g a a c c t t c c a g a a g t g g g c a g c t g t g G i g g t g c c t t c t g g a g a a g a g c a  
g a g a t a c a c g t g c c a t g t g c a g c a c g a g g g g c t g c c g g a g c c c c t a c c c t g a g a t g g g a g c c g t c t t c c a g c c c  
a c c a t c c c c a t c g t g g g c a t c g t g c t g g c c t g g c t g t c c t g g c t g t c c t a g c t g t c c t a g g a g c t g t g g t g g c t g  
5 t t g t g a t g t g t a g g a g g a a g a g c t c a g g t g g a a a g g a g g g a g c t g c t c t c a g g c t g c g t c c a g c a a c a g t g c c c a  
g g g c t c t g a t g a g t c t c t c a t c g c t t g t a a (SEQ ID NO:103);

Cw\*1404 :

g c t c c c a c t c c a t g a g g t a t t t c t C c a c a t c c g t g t c c c g g c c c g g c c g g g a g c c c c g c t t c a t c g c a g t g g g  
c t a c g t g g a c g a c a c g c a g t t c g t g c g g t t c g a c a g c g a c g c c g c g a g t c c g a g a g g g a g c c g c g g g c g c c g t g g  
10 g t g g a g c a g g a g g g c c g g a g t a t t g g g a c c g g g a g a c a c a g a a g t a c a a g c g c c a g g c a c a g g c t g a c c g a g t g a  
A c c t g c g g a a c c t g c g c g g c t a c t a c a a c c a g a g c g a g g c c g g g t c t c a c a c c c t c c a g t g g a t g t T t g g c t g c g a  
c c t g g g g c c c g a c g g g c g c c t c c t c c g c g g g t a t g a c c a g t C c g c c t a c g a c g g c a a g g a t t a c a t c g c c c t g a a c  
g a g g a T c t g c g c t c c t g g a c c g c c g g a c a c g g c g g c t c a g a t c a c c c a g c g c a a g t g g g a g g c g g c c c g t g a g g  
c g g a g c a g c g g a g a g c c t a c c t g g a g g g c a c g t g c g t g g a g t g g c t c c g c a g a t a c c t g g a g a a c g g g a a g g a g a c  
15 g c t g c a g c g c g c g g (SEQ ID NO:104);

Cw\*1405 :

g c t c c c a c t c c a t g a g g t a t t t c t C c a c a t c c g t g t c c c g g c c c g g c c g g g a g c c c c g c t t c a t c g c a g t g g g  
c t a c g t g g a c g a c a c g c a g t t c g t g c g g t t c g a c a g c g a c g c c g c g a g t c c g a g a g g g a g c c g c g g g c g c c g t g g  
g t g g a g c a g g a g g g c c g g a g t a t t g g g a c c g g g a g a c a c a g a a g t a c a a g c g c c a g g c a c a g a c t g a c c g a g t g a  
20 g c c t g c g g a a c c t g c g c g g c t a c t a c a a c c a g a g c g a g g c c g g g t c t c a c a c c c t c c a g t g g a t g t A t g g c t g c g a  
c c t g g g g c c c g a c g g g c g c c t c c t c c g c g g g t a t g a c c a g t C c g c c t a c g a c g g c a a g g a t t a c a t c g c c c t g a a c  
g a g g a T c t g c g c t c c t g g a c c g c c g g a c a c g g c g g c t c a g a t c a c c c a g c g c a a g t g g g a g g c g g c c c g t g a g g  
c g g a g c a g c g g a g a g c c t a c c t g g a g g g c a c g t g c g t g g a g t g g c t c c g c a g a t a c c t g g a g a a c g g g a a g g a g a c  
g c t g c a g c g c g c g g (SEQ ID NO:105);

25 Cw\*150201 :

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Cw\*150202 :

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Cw\*1503 :

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25 accatccccatcgtaggcatcgctgctggcctggctgtccctggctgtccctagctgtccctaggagctgtgAtggctg  
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5 gaggcaggaggggcccggaglattgggaccgggagacacagaactacaagcggcaggcacagactgaccgagtgaacc  
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15 ttgtgatgttaggaggaagagctcaggltggaaaaggaggagctgctctcaggctgcgtccagcaacagtgccca  
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20 cgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagttccaagaggggagccgcgggcgccgtgggtg  
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5 gggcctctgatgagtcctcctcgtttaa (SEQ ID NO:111);

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15 gcagcgcgcggaacacccaaagacacacgtgaccaccatcccgtctctgaccatgaggccacctgagggtgtgtg  
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25 gccgtcggaacctgcgcggctactacaaccagagcgaggccgggtctcacatcatccagaggatgtatggctgcga  
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Cw\*1508 :

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15 gtggagcaggaggggcccggagTatTgggaccgggagacacagaactacaagcgccaggcacagacTgaccgagTga  
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20 gctgcagcgcgcgg(SEQ ID NO:115);

Cw\*1510 :

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Cw\*1511:

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5 ctacTggacgacacgcagTtcgtgcggttcgacagcgacgccgcgagTccaagaggggagccgcgggcgcgcTgg  
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Cw\*1601:

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15 cgtggacgacacgcagTtcgtgcggttcgacagcgacgccgcgagTccaagaggggagccgcgggcgcgcTgggtg  
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15 ttgtTatgtgtaggaggaagagctcaggtggaaaaggagggagctgctctcaggctgcgtccagcaacagtgccca  
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Cw#160401 :

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20 cgtggacgacacgcagttcgtgcgggtcgacagcgacgccgagtcgaagaggggagccgcccggcgccgtgggtg  
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5 Cw\*1701 :

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10 tgcggaaactgcgcggctactacaaccagagcgaggccggttctcacaccatccagaggatgtatggctgcgacct  
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gacctgcgctcctggaccgcggcgacacggcggctcagatctccagcgcaagttggaggcgcccgtagggcgg  
agcagctgagagcctaccitggagggcgagtcgctggagtggctccgcggataccitggagaacgggaaggagacgt  
gcagcgcggaacgccccaaagacacagtgaccaccatcccgctcttgacctagggccaccttaggtgctgg  
15 gcccitgggttcttaccttgcggagatcacactgacctggcagcgggatggggaggaccaaactcaggacaccgagc  
ttigtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgttggtggccttctggacaagaaca  
gagatacacgtgccatgtgcagcacgaggggtgcaggagccctgcacctgagatggaagccgtcttccagccc  
accatcccaacttgggcatcgttcttggcccagctgtcctggcgtgcttggcgtgcttggcgtgcttagctgtcc  
taggagctgttggtgctgtgtgataC (SEQ ID NO:121);

20 Cw\*1702 :

atgcgggcatggcgccccgaaccttcatcctgctgcttccgggagccctggccctgatcgagacctgggcccgt  
cccactccatgaggtattctacaccgccgtgtccggcccgccggagagccccgcttcatcgcatgggctta  
cgtggacgacacgcagttcgtgcggttcgacagcgacgccgcgagtcgagaggggagccgcggcgccgtgggtg  
gagcaggagggggccggagtattgggaccgggagacacagaagtacaagcgccaggcacaggctgaccgagtgaaac

25 tgcggaaactgcgcggctactacaaccagagcgaggccggttctcacaccatccagaggatgtatggctgcgacct  
ggggcccgacggcgccctcctccggggtataaccagttcgcttacgacggcaaggattacatcgccctgaacgag  
gacctgcgctcctggaccgcggcgacacggcggctcagatctccagcgcaagttggaggcgcccgtagggcgg

agcagc t g a g a g c c t a c c t g g a g g g c g a g t g c g t g g a g t g g c t c c g c g g a t a c c t g g a g a a c g g g a a g g a g a c g c t  
g c a g c g c g c g g a a c g c c c a a a g a c a c a c g t g a c c c a c c a t c c c g t c t c t g a c c a t g a g g c c a c c c t g a g g t g c t g g  
g c c c t g g g c t t c t a c c c t g c g g a g a t c a c a c t g a c c t g g c a g c g g g a t g g g g a g g a c c a a a c t c a g g a c a c c g a g c  
t t g t g g a g a c c a g g c c a g c a g g a g a t g g a a c c t t c c a g a a g t g g g c a g c t g t g g t g g t g c c t t c t g g a c a a g a a c a  
5 g a g a t a c a c g t g c c a t g t g c a g c a c g a g g g g c t g c a g g a g c c c t g c a c c c t g a g a t g g a (SEQ ID NO:122) ;  
Cw\*1703 :  
a t g c g g g t c a t g g c g c c c c a a g c c c t c c t c c t g c t g c t c t c g g g a g c c c t g g c c c t g a t c g a g a c c t g g A c c g g c t  
c c c a c t c c a t g a g g t a t t t c t a c a c c g c c g t g t c c c g g c c c g g c g c g g a g a g c c c c g c t t c a t c g c a g t g g g c t a  
c g t g g a c g a c a c g c a g t t c g t g c g g t t c g a c a g c g a c g c c g c g a g t c c g a g a g g g g a g c c g c g g g c g c c g t g g g t g  
10 g a g c a g g a g g g g c c g g a g t a t t g g g a c c g g g a g a c a c a g a a g t a c a a g c g c c a g g c a c a g g c t g a c c g a g t g a a c c  
t g c g g a a a c t g c g c g g c t a c t a c a a c c a g a g c g a g g c c g g t t c t c a c a c c a t c c a g a g a t g t a t g g c t g c g a c c t  
g g g g c c c g a c g g g c g c c t c c t c c g c g g g t a t a a c c a g t t c g c c t a c g a c g g c a a g g a t t a c a t c g c c c t g a a c g a g  
g a c c t g c g c t c c t g g a c c g c g g c g g a c a c g g c g g c t c a g a t c t c c c a g c g c a a g t t g g a g g c g g c c c g t g a g g c g g  
a g c a g c t g a g a g c c t a c c t g g a g g g c g a g t g c g t g g a g t g g c t c c g c g g a t a c c t g g a g a a c g g g a a g g a g a c g c t  
15 g c a g c g c g c g g a a c g c c c a a a g a c a c a c g t g a c c c a c c a t c c c g t c t c t g a c c a t g a g g c c a c c c t g a g g t g c t g g  
g c c c t g g g c t t c t a c c c t g c g g a g a t c a c a c t g a c c t g g c a g c g g g a t g g g g a g g a c c a a a c t c a g g a c a c c g a g c  
t t g t g g a g a c c a g g c c a g c a g g a g a t g g a a c c t t c c a g a a g t g g g c a g c t g t g g t g g t g c c t t c t g g a c a a g a a c a  
g a g a t a c a c g t g c c a t g t g c a g c a c g a g g g g c t g c a g g a g c c c t g c a c c c t g a g a t g g a a g c c g t c t t c c a g c c c  
a c c a t c c c c a a c t t g g g c a t c g t t t c t g g c c c a g c t g t c c t g g c t g t c c t g g c t g t c c t a g c t g t c c  
20 t a g g a g c t g t g g t c g c t g c t g t g a t a c (SEQ ID NO:123) ;  
Cw\*1801 :  
a t g c g g g t c a t g g c g c c c c g a g c c c t c c t c c t g c t g c t c t c g g g a g g c c t g g c c c t g a c c g a g a c c t g g g c c t g c t  
c c c a c t c c a t g a g g t a t t t c g a c a c c g c c g t g t c c c g g c c c g g c g c g g a g a g c c c c g c t t c a t c t c a g t g g g c t a  
c g t g g a c g a c a c g c a g t t c g t g c g g t t c g a c a g c g a c g c c g c g a g t c c g a g a g g g g a g c c c c g g g c g c c g t g g g t g  
25 g a g c a g g a g g g g c c g g a g t a t t g g g a c c g g g a g a c a c a g a a g t a c a a g c g c c a g g c a c a g g c t g a c c g a g t g a a c c  
t g c g g a a a c t g c g c g g c t a c t a c a a c c a g a g c g a g g a c g g g t c t c a c a c c c t c c a g a g g a t g t t t g g c t g c g a c c t  
g g g g c c g g a c g g g c g c c t c c t c c g c g g g t a t a a c c a g t t c g c c t a c g a c g g c a a g g a t t a c a t c g c c c t g a a c g a g

- gatctgcgtccctggaccgccgacacggcggctcagatcaccagcgcaagtgggaggcgcccgtagggcgg  
 agcagcggagagcctacctggagggcacgtgcgtggagtggctccgcagataacctggagaacgggaaggagacgct  
 gcagcgcgcggaacacccaaagacacacgtgaccacacatcccgtctctgaccaatgaggccacccatgaggctgctgg  
 gcccctgggcttctacctgctggagatcacactgacctggcagtgggatggggaggaccacaaactcaggacaccgagc  
 5 ttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtggtggcttcttggagaagagca  
 gagatacacgtgccatgtgcagcacgaggggctgccggagccccctacccctgagatggAagccgtcttcccagccc  
 accatccccatcgtgggcatcgttgcctggcctggctgtcttggTtgtcttagctgtcttaggagctgtggtggctg  
 ttgtgatgtgtaggaggaagagctcagggtgaaaaggaggagctgtctcaggctgcgtccagcaacagtgtccca  
 gggctctgatgagctctctcatcgttgtaa (SEQ ID NO:124);
- 10 Cw\*1802 :
- atgcgggtcatggcgccccgagccctctctctgtctgtctcgggaggccctggccctgaccgagaccctgggcttgc  
 cccactccatgaggatatttcgacaccgccgtgtcccgcccgcccgaggagccccgttcatctcagtgggctta  
 cgtggacgacacgcagttcgtgcggttcgacagcgacgcccgagctccgagaggggagccCcgggcgccgtgggtg  
 gaggcaggagggggcgaglatgggaccgggagacacagaagtacaagcgccaggcacaggctgaccgagtgaacc  
 15 tgcggaaactgcgcggctactacaaccagagcaggacgggctcacacccctccagaggatgtttggctgcgacct  
 ggggcccggacggcgccctctcgcgggtataaccagttccctacgacggcaaggatlacatcggcctgaacgag  
 gaTctgcgtccctggaccgccgacacggcggctcagatcaccagcgcaagtgggaggcgcccgtagggcgg  
 agcagcggagagcctacctggagggcacgtgcgtggagtggctccgcagataacctggagaacgggaaggagacgct  
 gcagcgcgcggaacacccaaagacacacgtgaccacacatcccgtctctgaccaatgaggccacccatgaggctgctgg  
 20 gcccctgggcttctacctgctggagatcacactgacctggcagtgggatggggaggaccacaaactcaggacaccgagc  
 ttgtggagaccaggccagcaggagatggaaccttccagaagtgggcagctgtggtggcttcttggagaagagca  
 gagatacacgtgccatgtgcagcacgaggggctgccggagccccctacccctgagatggAagccgtcttcccagccc  
 accatccccatcgtgggcatcgttgcctggcctggctgtcttggctgtcttagctgtcttaggagctgtggtggctg  
 ttgtgatgtgtaggaggaagagctcagggtgaaaaggaggagctgtctcaggctgcgtccagcaacagtgtccca  
 25 gggctctgatgagctctctcatcgttgtaa (SEQ ID NO:125);

In the following, Probe Lists C1 and C2 are

shown In Tables 9-1 to 9-4 and Tables 10-1 to 10-4 respectively.

Table 9-1

Probe No.	Base Sequence
0	c acc ctc cag tgg atg tG ( SEQ ID No: 1 2 6)
1	c cgc ggg tat gac cag tA ( SEQ ID No: 1 2 7)
2	g acc gcc gcg gac acC ( SEQ ID No: 1 2 8)
3	ag aag tgg gca gct gtg A ( SEQ ID No: 1 2 9)
4	c ctc ctc cgc ggg tat A ( SEQ ID No: 1 3 0)
5	g cgc tcc tgg acc gcT ( SEQ ID No: 1 3 1)
6	g cac gag ggg ctg ccA ( SEQ ID No: 1 3 2)
7	ct gtc cta gga gct gtg A ( SEQ ID No: 1 3 3)
8	c acc ctc cag agg atg tC ( SEQ ID No: 1 3 4)
9	gg gag gcg gcc cgt gT ( SEQ ID No: 1 3 5)
10	ggg cgc ctc ctc cgc A ( SEQ ID No: 1 3 6)
11	c aag tgg gag gcg gcc T ( SEQ ID No: 1 3 7)
12	c cgt gag gcg gag cag T ( SEQ ID No: 1 3 8)
13	a gtg aac ctg cgg aaa ctA ( SEQ ID No: 1 3 9)
14	cc ctg ggc ttc tac cct A ( SEQ ID No: 1 4 0)
15	g acc gcc gcg gac acA ( SEQ ID No: 1 4 1)
16	gct gtg tcc cgg ccc A ( SEQ ID No: 1 4 2)
17	g acc gcc gcg gac acG ( SEQ ID No: 1 4 3)
18	cc ctg aga tgg gag ccA ( SEQ ID No: 1 4 4)
19	gg tct cac acc ctc cag A ( SEQ ID No: 1 4 5)
20	cgc ggg tat gac cag tC ( SEQ ID No: 1 4 6)
21	gcc tac ctg gag ggc gA ( SEQ ID No: 1 4 7)
22	c tcc cac tcc atg agg tG ( SEQ ID No: 1 4 8)
23	cgc ggg cat gac cag ttA ( SEQ ID No: 1 4 9)
24	g gac caa act cag gac acT ( SEQ ID No: 1 5 0)
25	c aac cag agc gag gcc A ( SEQ ID No: 1 5 1)
26	ag gcc agg tct cac atc A ( SEQ ID No: 1 5 2)
27	g aag tgg gca gct gtg G ( SEQ ID No: 1 5 3)
28	gcg gac acg gcg gcC ( SEQ ID No: 1 5 4)
29	at ggc tgc gac gtg ggA ( SEQ ID No: 1 5 5)
30	g gcc ggg tct cac atc A ( SEQ ID No: 1 5 6)

Table 9-2

Probe No.	Base Sequence
31	c atc atc cag agg atg taC ( SEQ ID No: 157)
32	c cgc aga tac ctg aag aaT ( SEQ ID No: 158)
33	ct cac acc ctc cag agC ( SEQ ID No: 159)
34	ctc ctc cgc ggg tat gT ( SEQ ID No: 160)
35	ca cag act gac cga gtg aA ( SEQ ID No: 161)
36	cga gtg aac ctg cgg aaA ( SEQ ID No: 162)
37	gg atg tat ggc tgc gac G ( SEQ ID No: 163)
38	gcc tac ctg gag ggc cT ( SEQ ID No: 164)
39	gac cgg gag aca cag aaC ( SEQ ID No: 165)
40	g gag ccc cac ttc atc G ( SEQ ID No: 166)
41	cga gtg agc ctg cgg aaA ( SEQ ID No: 167)
42	cgc ggg tat gac cag ttA ( SEQ ID No: 168)
43	g gag gcg gcc cgt gC ( SEQ ID No: 169)
44	c tac aac cag agc gag gA ( SEQ ID No: 170)
45	cgt gag gcg gag cag cT ( SEQ ID No: 171)
46	cta gct gtc cta gga gct A ( SEQ ID No: 172)
47	ggc tac gtg gac gac acA ( SEQ ID No: 173)
48	gc cgc gga gag ccc cA ( SEQ ID No: 174)
49	g aga tac acg tgc cat gtT ( SEQ ID No: 175)
50	ga ggg gag ccg cgg gA ( SEQ ID No: 176)
51	c atc gca gtg ggc tac C ( SEQ ID No: 177)
52	c tgc gac ctg ggg ccG ( SEQ ID No: 178)
53	tc tcc aca tcc gtg tcc T ( SEQ ID No: 179)
54	c aag cgc cag gca cag G ( SEQ ID No: 180)
55	gg acc gcc gcg gac aA ( SEQ ID No: 181)
56	ctc acc ctg aga tgg gG ( SEQ ID No: 182)
57	tg tgc gtg gag tgg ctG ( SEQ ID No: 183)
58	cc atc tct gac cat gag gT ( SEQ ID No: 184)
59	ac ctg gag aac ggg aag A ( SEQ ID No: 185)
60	c cgc ggg tat aac cag tT ( SEQ ID No: 186)

Table 9-3

Probe No.	Base Sequence
61	g gag ccg cgg gcg cG ( SEQ ID No: 187)
62	t ccg aga ggg gag ccC ( SEQ ID No: 188)
63	g agg tat ttc tac acc gcT ( SEQ ID No: 189)
64	c gac gcc gcg agt ccA ( SEQ ID No: 190)
65	gt cca aga ggg gag ccC ( SEQ ID No: 191)
66	gcg ccg tgg gtg gag A ( SEQ ID No: 192)
67	c acc ctc cag agg atg tA ( SEQ ID No: 193)
68	g atc acc cag cgc aag tT ( SEQ ID No: 194)
69	g acg ctg cag cgc gcA ( SEQ ID No: 195)
70	c tct gat gag tct ctc atc A ( SEQ ID No: 196)
71	gag cca tct tcc cag ccT ( SEQ ID No: 197)
72	ga gcc tac ctg gag ggA ( SEQ ID No: 198)
73	t gcg gcg gag cag gaC ( SEQ ID No: 199)
74	aac ctg cgc ggc tac taT ( SEQ ID No: 200)
75	g tct cac acc ctc cag aaT ( SEQ ID No: 201)
76	a gct gtg gtc acc gct aA ( SEQ ID No: 202)
77	c acc ctc cag agg atg tT ( SEQ ID No: 203)
78	ag gac ggg tct cac atc A ( SEQ ID No: 204)
79	ac atc atc cag agg atg tC ( SEQ ID No: 205)
80	tgc tct cag gct gcg tG ( SEQ ID No: 206)
81	c cgc ggg tat gac cag tT ( SEQ ID No: 207)
82	g gag acg ctg cag cgc A ( SEQ ID No: 208)
83	g ccc ctc acc ctg agC ( SEQ ID No: 209)
84	ggg agc tgc tct cag gT ( SEQ ID No: 210)
85	cgt acg gcg gag cag cT ( SEQ ID No: 211)
86	acc ctc cag agg atg taC ( SEQ ID No: 212)
87	tgg gag gcg gcc cgt A ( SEQ ID No: 213)
88	cgc aga tac ctg gag aac A ( SEQ ID No: 214)
89	gcc tac ctg gag ggc G ( SEQ ID No: 215)
90	ga tac ctg gag aac ggg G ( SEQ ID No: 216)

Table 9-4

Probe No.	Base Sequence
91	ac ctg cgc tcc tgg acT (SEQ ID No: 217)
92	g cgc tcc tgg acc gcG (SEQ ID No: 218)
93	a gag ccc cgc ttc atc G (SEQ ID No: 219)
94	c acc ctc cag tgg atg tA (SEQ ID No: 220)
95	cag tcc gcc tac gac gT (SEQ ID No: 221)
96	a cag gct gac cga gtg G (SEQ ID No: 222)
97	cac tcc atg agg tat ttc tC (SEQ ID No: 223)
98	c acc ctc cag tgg atg tT (SEQ ID No: 224)
99	a cag gct gac cga gtg aA (SEQ ID No: 225)
100	atc gcc ctg aac gag gaT (SEQ ID No: 226)
101	gc ctc ctc cgc ggg C (SEQ ID No: 227)
102	tc atg gcg ccc cga acT (SEQ ID No: 228)
103	cgc ggg cat gac cag tT (SEQ ID No: 229)
104	cgc ggg cat gac cag tC (SEQ ID No: 230)
105	gt gcg gcg gag cag cA (SEQ ID No: 231)
106	gct gtg gtg gct gtt gtT (SEQ ID No: 232)
107	cgt gcg gcg gag cag T (SEQ ID No: 233)
108	tg gtc gct gct gtg ata C (SEQ ID No: 234)
109	gg ctg cag gag ccc tG (SEQ ID No: 235)
110	cc ctg atc gag acc tgg A (SEQ ID No: 236)
111	cc ctc acc ctg aga tgg A (SEQ ID No: 237)
112	ggc ctg gct gtc ctg gT (SEQ ID No: 238)



Table 10-1

Probe No.

Base Sequence

0	g tgg atg tGt ggc tgc g (SEQ ID No: 239)
1	at gac cag tAc gcc tac g (SEQ ID No: 240)
2	gcg gac acC gcg gct c (SEQ ID No: 241)
3	gca gct gtg Atg gtg cct (SEQ ID No: 242)
4	cgc ggg tat Aac cag ttc (SEQ ID No: 243)
5	tgg acc gcT gcg gac ac (SEQ ID No: 244)
6	ggg ctg ccA gag ccc c (SEQ ID No: 245)
7	gga gct gtg Atg gct gtt (SEQ ID No: 246)
8	g agg atg tCt ggc tgc g (SEQ ID No: 247)
9	g gcc cgt gTg gcg gag (SEQ ID No: 248)
10	ctc ctc cgc Agg tat gac (SEQ ID No: 249)
11	g gcg gcc Tgt gag gcg (SEQ ID No: 250)
12	cg gag cag Tgg aga gcc (SEQ ID No: 251)
13	g cgg aaa ctA cgc ggc ta (SEQ ID No: 252)
14	ttc tac cct Acg gag atc a (SEQ ID No: 253)
15	gcg gac acA gcg gct c (SEQ ID No: 254)
16	c cgg ccc Agc cgc gg (SEQ ID No: 255)
17	gcg gac acG gcg gct c (SEQ ID No: 256)
18	a tgg gag ccA tct tcc ca (SEQ ID No: 257)
19	acc ctc cag Agg atg tat g (SEQ ID No: 258)
20	t gac cag tCc gcc tac g (SEQ ID No: 259)
21	g gag ggc gAg tgc gtg (SEQ ID No: 260)
22	cc atg agg tGt ttc tac ac (SEQ ID No: 261)
23	t gac cag ttA gcc tac gac (SEQ ID No: 262)
24	t cag gac acT gag ctt gtg (SEQ ID No: 263)
25	gc gag gcc Agg tct cac (SEQ ID No: 264)
26	tct cac atc Atc cag agg a (SEQ ID No: 265)
27	ca gct gtg Gtg gtg cct (SEQ ID No: 266)
28	acg gcg gcC cag atc ac (SEQ ID No: 267)
29	gac gtg ggA ccc gac g (SEQ ID No: 268)
30	g agg atg taC ggc tgc ga (SEQ ID No: 269)

Table 10-2

Probe No.	Base Sequence
31	c ctg aag aaT ggg aag gag ( SEQ ID No : 270 )
32	c ctc cag agC atg tac gg ( SEQ ID No : 271 )
33	gc ggg tat gTc cag tac g ( SEQ ID No : 272 )
34	c cga gtg aAc ctg cgg a ( SEQ ID No : 273 )
35	ctg cgg aaA ctg cgc gg ( SEQ ID No : 274 )
36	c tgc gac Gtg ggg ccc ( SEQ ID No : 275 )
37	g gag ggc cTg tgc gtg ( SEQ ID No : 276 )
38	g aca cag aaC tac aag cgc ( SEQ ID No : 277 )
39	cac ttc atc Gca gtg ggc ( SEQ ID No : 278 )
40	gcc cgt gCg gcg gag ( SEQ ID No : 279 )
41	g agc gag gAc ggg tct c ( SEQ ID No : 280 )
42	g gag cag cTg aga gcc t ( SEQ ID No : 281 )
43	cta gga gct Atg gtg gct ( SEQ ID No : 282 )
44	g gac gac acA cag ttc gt ( SEQ ID No : 283 )
45	ga gag ccc cAc ttc atc g ( SEQ ID No : 284 )
46	g tgc cat gTt cag cac ga ( SEQ ID No : 285 )
47	ccg cgg gAg ccg tgg ( SEQ ID No : 286 )
48	tg ggc tac Ctg gac gac ( SEQ ID No : 287 )
49	ctg ggg ccG gac ggg ( SEQ ID No : 288 )
50	c gtg tcc Tgg ccc ggc ( SEQ ID No : 289 )
51	ag gca cag Gct gac cga ( SEQ ID No : 290 )
52	c gcg gac aAg gcg gct ( SEQ ID No : 291 )
53	tg aga tgg gGg cca tct t ( SEQ ID No : 292 )
54	g gag tgg ctG cgc aga ta ( SEQ ID No : 293 )
55	ac cat gag gTc acc ctg a ( SEQ ID No : 294 )
56	aac ggg aag Aag acg ctg ( SEQ ID No : 295 )
57	at aac cag tTc gcc tac ga ( SEQ ID No : 296 )
58	cgg gcg cGg tgg gtg ( SEQ ID No : 297 )
59	ggg gag ccC cgg gcg ( SEQ ID No : 298 )
60	tac acc gcT gtg tcc cg ( SEQ ID No : 299 )

Table 10-3

Probe No.	Base Sequence
61	gcg agt ccA aga ggg ga ( SEQ ID No: 300)
62	gg gtg gag Aag gag ggg ( SEQ ID No: 301)
63	ag agg atg tAt ggc tgc g ( SEQ ID No: 302)
64	g cgc aag tTg gag gcg g ( SEQ ID No: 303)
65	cag cgc gcA gaa ccc c ( SEQ ID No: 304)
66	g gct gcg tGc agc aac a ( SEQ ID No: 305)
67	tcc cag ccT acc atc cc ( SEQ ID No: 306)
68	ctg gag ggA ctg tgc gt ( SEQ ID No: 307)
69	g gag cag gaC aga gcc ta ( SEQ ID No: 308)
70	c ggc tac taT aac cag agc ( SEQ ID No: 309)
71	c ctc cag aaT atg tat ggc ( SEQ ID No: 310)
72	tc acc gct aAg atg tgt ag ( SEQ ID No: 311)
73	ag agg atg tTt ggc tgc g ( SEQ ID No: 312)
74	at gac cag tTc gcc tac g ( SEQ ID No: 313)
75	ggg ctg caA gag ccc c ( SEQ ID No: 314)
76	gc tct cag gTt gcg tgc a ( SEQ ID No: 315)
77	g gcc cgt Acg gcg gag ( SEQ ID No: 316)
78	ctg gag aac Agg aag aag a ( SEQ ID No: 317)
79	g gag ggc Gcg tgc gtg ( SEQ ID No: 318)
80	c ctc cag agC atg tat gg ( SEQ ID No: 319)
81	gag aac ggg Gag aag acg ( SEQ ID No: 320)
82	tcc tgg acT gcc gcg g ( SEQ ID No: 321)
83	tgg acc gcG gcg gac a ( SEQ ID No: 322)
84	gc ttc atc Gca gtg ggc ( SEQ ID No: 323)
85	ag tgg atg tAt ggc tgc g ( SEQ ID No: 324)
86	cc tac gac gTc aag gat ta ( SEQ ID No: 325)
87	c cga gtg Ggc ctg cgg ( SEQ ID No: 326)
88	gg tat ttc tCc aca tcc gt ( SEQ ID No: 327)
89	ag tgg atg tTt ggc tgc g ( SEQ ID No: 328)
90	g aac gag gaT ctg cgc tc ( SEQ ID No: 329)

Table 10-4

Probe No.	Base Sequence
91	c cgc ggg Cat gac cag ( SEQ ID No: 3 3 0)
92	ccc cga acT ctc ctc ct ( SEQ ID No: 3 3 1)
93	c cgc ggg Cat gac cag ( SEQ ID No: 3 3 2)
94	g gag cag cAg aga gcc t ( SEQ ID No: 3 3 3)
95	g gct gtt gtT atg tgt agg ( SEQ ID No: 3 3 4)
96	t gtg gtc gcT gct gtg at ( SEQ ID No: 3 3 5)
97	g gag ccc tGc acc ctg ( SEQ ID No: 3 3 6)
98	g acc tgg Acc ggc tcc ( SEQ ID No: 3 3 7)
99	ctg aga tgg Aag ccg tct ( SEQ ID No: 3 3 8)
100	ct gtc ctg gTt gtc cta g ( SEQ ID No: 3 3 9)

Table 11-1

Allele Number	Probe Number for Detection					
Cw*0102	0	1	2	3		
Cw*0103	4					
Cw*0104	5	6	7			
Cw*0105	8					
Cw*0106	9					
Cw*0107	10					
Cw*0108	11					
Cw*0109	12					
Cw*020201	13					
Cw*020202	14					
Cw*020203	15	12				
Cw*020204	16	17	18			
Cw*020205	16	19	20	17	12	21
Cw*0203	9	21				
Cw*0204	22					
Cw*0205	16	20	17	12	21	
Cw*0206	23	21				
Cw*030201	24	18				
Cw*030202	20	24				
Cw*030301	25	26	27			
Cw*030302	28					
Cw*030303	29					
Cw*030401	30	24				
Cw*030402	30	31	32			
Cw*0305	33	32				
Cw*0306	34					
Cw*0307	35	36	30	37	38	32
Cw*0308	39	30	24			
Cw*0309	40	30	38	32		
Cw*0310	41	30	37	38	32	
Cw*0311	25	26				

Table 11-2

Allele Number	Probe Number for Detection			
Cw*0312	25	42		
Cw*0313	25	27		
Cw*0314	43	32		
Cw*0315	44	20	38	32
Cw*0316	37	20	17	45
Cw*040101	46			
Cw*040102	47			
Cw*0403	48	49		
Cw*0404	50	45		
Cw*0405	51			
Cw*0406	48	52	45	
Cw*0407	53	54		
Cw*0408	50	38		
Cw*0410	50			
Cw*0501	36	55	56	
Cw*0502	57			
Cw*0503	58			
Cw*0504	20	55	59	
Cw*0505	37	60	55	59
Cw*0506	61			
Cw*0602	62	12	7	
Cw*0603	63	62	20	12
Cw*0604	62	45		
Cw*0605	64	65	20	17
Cw*0606	62	7		
Cw*0607	66			
Cw*0608	44	20	17	12 21
Cw*0609	62	60	12	
Cw*070101	67	68	69	70
Cw*070102	71			

Table 11-3

Allele Number		Probe Number for Detection				
Cw*070201	8	68	70			
Cw*0703	72					
Cw*070401	73	70				
Cw*070402	74					
Cw*0705	75					
Cw*0706	76					
Cw*0707	36	67	20	68	69	
Cw*0708	77	20	68	69		
Cw*0709	36	44	67	20	68	69
Cw*0710	78	79	20	68	69	
Cw*0711	73	80				
Cw*0712	73					
Cw*0713	8	81	68	69		
Cw*0714	82					
Cw*0715	8	21	69			
Cw*0716	39	67	20	68	69	
Cw*0717	8	83				
Cw*0718	84					
Cw*080101	85	56				
Cw*080102	86	60	87			
Cw*0802	55	56				
Cw*0803	88	7				
Cw*0804	55	45	59			
Cw*0805	54	60	55	59		
Cw*0806	89	88				
Cw*0807	55	68	59			
Cw*0808	33	59				
Cw*0809	90					
Cw*120201	86	5	7			
Cw*120202	86	5	6	7		
Cw*120203	67	5				
Cw*120301	54	91	7			

Table 11-4

Allele Number		Probe Number for Detection					
Cw*120302	92	12					
Cw*120401	93	54	36	94	20	17	12
Cw*120402	54	36	91	7			
Cw*1205	36	91	7				
Cw*1206	95						
Cw*1207	96						
Cw*1208	39	86	5	6	7		
Cw*140201	97	20	27				
Cw*140202	97	98	20				
Cw*1403	97	64	20	27			
Cw*1404	97	99	98	20	100		
Cw*1405	97	94	20	100			
Cw*150201	23	7					
Cw*150202	48	39	36	101	23	45	
Cw*1503	54	23	7				
Cw*1504	20	45	7				
Cw*150501	102						
Cw*150502	101	103	7				
Cw*1506	101	7					
Cw*1507	48	39	101	23	45		
Cw*1508	48	39	36	30	101	23	
Cw*1509	101	104	45				
Cw*1510	39	36	101	23	45		
Cw*1511	16	48	36	101	23	45	
Cw*1601	105	106					
Cw*1602	36	105	106				
Cw*160401	107	106					
Cw*1701	108						
Cw*1702	109						
Cw*1703	110						
Cw*1801	111	112					
Cw*1802	62	100	111				



Table 12-1

Allele Number	Probe Number for Detection					
Cw*0102	0	1	2	3		
Cw*0103	4					
Cw*0104	5	6	7			
Cw*0105	8					
Cw*0106	9					
Cw*0107	10					
Cw*0108	11					
Cw*0109	12					
Cw*020201	13					
Cw*020202	14					
Cw*020203	15	12				
Cw*020204	16	17	18			
Cw*020205	16	19	20	17	12	21
Cw*0203	9	21				
Cw*0204	22					
Cw*0205	16	20	17	12	21	
Cw*0206	23	21				
Cw*030201	24	18				
Cw*030202	20	24				
Cw*030301	25	26	27			
Cw*030302	28					
Cw*030303	29					
Cw*030401	26	24				
Cw*030402	26	30	31			
Cw*0305	32	31				
Cw*0306	33					
Cw*0307	34	35	26	36	37	31
Cw*0308	38	26	24			
Cw*0309	39	26	37	31		
Cw*0310	35	26	36	37	31	
Cw*0311	25	26				

Table 12-2

Allele Number	Probe Number for Detection				
Cw*0312	25	23			
Cw*0313	25	27			
Cw*0314	40	31			
Cw*0315	41	20	37	31	
Cw*0316	36	20	17	42	
Cw*040101	43				
Cw*040102	44				
Cw*0403	45	46			
Cw*0404	47	42			
Cw*0405	48				
Cw*0406	45	49	42		
Cw*0407	50	51			
Cw*0408	47	37			
Cw*0410	47				
Cw*0501	35	52	53		
Cw*0502	54				
Cw*0503	55				
Cw*0504	20	52	56		
Cw*0505	36	57	52	56	
Cw*0506	58				
Cw*0602	59	12	7		
Cw*0603	60	59	20	12	
Cw*0604	59	42			
Cw*0605	61	59	20	17	
Cw*0606	59	7			
Cw*0607	62				
Cw*0608	41	20	17	12	21
Cw*0609	59	57	12		
Cw*070101	63	64	65	66	
Cw*070102	67				
Cw*070201	8	64	66		

Table 12-3

Allele Number		Probe Number for Detection			
Cw*0703	68				
Cw*070401	69	66			
Cw*070402	70				
Cw*0705	71				
Cw*0706	72				
Cw*0707	38	35	40	42	
Cw*0708	73	40	42		
Cw*0709	38	35	41	40	42
Cw*0710	26	8	20	64	42
Cw*0711	69	66			
Cw*0712	69				
Cw*0713	8	74	64	42	
Cw*0714	30	64	40	42	
Cw*0715	8	21			
Cw*0716	38	40	42		
Cw*0717	8	75			
Cw*0718	76				
Cw*080101	42	53			
Cw*080102	30	57	77		
Cw*0802	52	53			
Cw*0803	78	7			
Cw*0804	52	42	56		
Cw*0805	51	57	52	56	
Cw*0806	79	78			
Cw*0807	52	64	56		
Cw*0808	80	56			
Cw*0809	81				
Cw*120201	30	5	7		
Cw*120202	30	5	6	7	
Cw*120203	63	5			
Cw*120301	51	82	7		

Table 12-4

Allele Number	Probe Number for Detection						
Cw*120302	83	12					
Cw*120401	84	51	35	85	20	17	12
Cw*120402	51	35	82	7			
Cw*1205	35	82	7				
Cw*1206	86						
Cw*1207	87						
Cw*1208	38	30	5	6	7		
Cw*140201	88	20	27				
Cw*140202	88	89	20				
Cw*1403	88	61	20	27			
Cw*1404	88	34	89	20	90		
Cw*1405	88	85	20	90			
Cw*150201	23	7					
Cw*150202	45	38	35	91	23	42	
Cw*1503	51	23	7				
Cw*1504	20	42	7				
Cw*150501	92						
Cw*150502	91	74	7				
Cw*1506	91	7					
Cw*1507	45	38	91	23	42		
Cw*1508	45	38	35	26	91	23	
Cw*1509	91	20	42				
Cw*1510	38	35	91	23	42		
Cw*1511	16	45	35	91	23	42	
Cw*1601	94	95					
Cw*1602	35	94	95				
Cw*160401	12	95					
Cw*1701	96						
Cw*1702	97						
Cw*1703	98						
Cw*1801	99	100					
Cw*1802	59	90	99				

(Example 7)

Probes for identification of HLA-DP allele

Extraction of DNA from 1 ml of human blood was performed using GFX Genomic Blood DNA Purification  
5 Kit from Amersham Biosciences in the same manner as in Example 1.

Next, quantitative PCR was carried out in the same manner as in Example 1 except that probes in the probe list 1 in Tables 13-1 to 13-3 or 14-1 to 14-3  
10 were used and 3  $\mu$ l of the mixed primers contains 1  $\mu$ l of respective solutions of the following primers (10 pmol/ $\mu$ l):

AAACACGGTCACCTCAGGGGGAT (SEQ ID NO: 245)

GGCCTGAGTGTGGTTGGAACG (SEQ ID NO: 246)

15 CCAGCTCGTAGTTGTGTCTGCA (SEQ ID NO: 247)

After PCR amplification, referring to Amp Plot and Dissociation curves on a display of 5700 software, and to the list in Table 15-1 for the probes in Table 13-1, or to the list in Tables 15-2 to 15-5 for the  
20 probes in Tables 13-2 to 13-3, it was identified as DPA1\*010301 and DPB1\*0901.

(Example 8)

Extraction of DNA from 1 ml of human blood was performed in the same way as in Example 1. PCR of  
25 human HLA-DP was then performed in the same manner as in Example 2 except that 6  $\mu$ l of the mixed primer consisting of 1  $\mu$ l each of the solutions containing

the following sequences at 10 pmol/ $\mu$ l respectively and 9  $\mu$ l of ultra pure water.

AAACACGGTCACCTCAGGGGGAT (SEQ ID NO: 245)

GGCCTGAGTGTGGTTGGAACG (SEQ ID NO: 246)

5 CCAGCTCGTAGTTGTGTCTGCA (SEQ ID NO: 247)

CCATGTGTCAACTTATGCC (SEQ ID NO: 248)

AGAATTACCTTTTCCAG (SEQ ID NO: 250)

AGAATTACGTTTTCCAG (SEQ ID NO: 251)

At the same time, a DNA microarray was prepared  
10 to identify the allele in the specimen described  
above in the same manner as in Example 2, except that  
probes in Tables 14-1 and 14-2 were used to form the  
probe spots respectively.

Then, hybridization was performed using the  
15 above specimen and the prepared DNA microarray in the  
same manner as in Example 2. Fluorometry measurement  
was conducted with GenePix4000B (Axon). Referring to  
the list in Table 16-1 when the probes in Table 14-1  
were used, or to the list in Tables 16-2 to 16-5 when  
20 the probes in Table 14-2 were used, the sample was  
identified as DPA1\*010301 and DPB1\*0901.

#### Allele list

DPA1\*010301 :

25 ccatgtgtcaacttatgccgcgtttgtacagacgcatagaccaacaggGgagtttatgtttgaatttgatgaAgat  
gagatgtttcctatgttggaatctggacaagaaggagaccgtctggcatctggaggagtttggccAagccttttcccttg  
aggctcagggcgggctggctaacattgctatatatgaacaacaacttgaalaccttgatccagcgttccaaccacac

475

tcaggccaccaac (SEQ ID NO: 1) ;

DPA1\*010302 :

gcgtttgtacagacgcatagaccaacaggAgagtttattgtttgaatttgatgaagatgagatgttctatgtggatc  
tggacaagaaggagaccgtctggcatctggaggagtttggccaagccttttccctttgaggctcagggcgggctggc

5 taacattgctatattgaacaacaacttgaataccttgatccagcgttccaaccacactcaggccaccaac (SEQ  
ID NO: 2) ;

DPA1\*0104 :

gccgcgtttgtacagacgcatagaccaacaggggagtttattgtttgaatttgatgaCgatgagatgttctatgtgg  
atctggacaagaaggagaccgtctggcatctggaggagtttggccaagccttttccctttgaggctcagggcgggct

10 ggctaacattgctatattgaacaacaacttgaataccttgatccagcgttccaaccacactcaggccaccaac (SE  
Q ID NO: 3) ;

DPA1\*0105 :

gccgcgtttgtacagacgcatagaccaacaggggagtttattgtttgaatttgatgaagatgagatgttctatgtgg  
atctggacaagaaggagaccgtctggcatctggaggagtttggccaagccttttccctttgaggctcagggcgggct

15 ggctaacattgctatattgaacaacaacttgaataccttgatccagcgttccaaccacactcaggccgccaAT (SE  
Q ID NO: 4) ;

DPA1\*0106 :

ccatgtgtcaacttattgccgcgtttgtacagacgcatagaccaacaggggagtttattgtttgaatttgatgaagat  
gagcagttctatgtggatctggataaAaaggagaccgtctggcatctggaggagtttggccaagccttttccctttg

20 aggcctcagggcgggctggctaacattgctatattgaacaacaacttgaataccttgatccagcgttccaaccacac  
tcaggccaccaac (SEQ ID NO: 5) ;

DPA1\*0107 :

catgtgtcaacttattgccgcgtttgtacagacgcatagaccaacaggggagtttattgtttgaatttgatgaagatg  
agatgttctatgtggatctggacaagaaggagaccgtctggcatctggaggagtttggccaaAccttttccctttga

25 ggctcagggcgggctggctaacattgctatattgaacaacaacttgaataccttgatccagcgttccaaccacact  
caggccaccaac (SEQ ID NO: 6) ;

DPA1\*0108 :

ccaatgltgcaacttatgccgcgtttgtacagacgcatagaccaacaggggagtttatgtttgaatttgatgaCgat  
gagaigtcttatgttgatctggacaagaaggagaccgtctggcatctggaggagtttggccGagccttttcccttg  
aggctcagggcgggctggctaacaattgctatattgaacaacaacttgaataccttgatccagcgttccaaccacac  
tcaggccaccaac (SEQ ID NO: 7) ;

5 DPA1\*020101 :

ccaatgltgcaacttatgccgcgtttgtacagacCcatagaccaacaggggagtttatgtttgaatttgatgaagat  
gagcagttcttatgttgatctggataaAaaggagaccgtctggcatctggaggagtttggccgagccttttcccttg  
aggctcagggcgggctggctaacaattgctatattgaacaacaacttgaataccttgatccagcgttccaaccacac  
tcaggccgccaat (SEQ ID NO: 8) ;

10 DPA1\*020102 :

ccaatgltgcaacttatgccgcgtttgtacagacgcatagaccaacaggggagtttatgtttgaatttgatgaagat  
gagcagttcttatgttgatctggataaAaaggagaccgtctggcatctggaggagtttggccgagccttttcccttg  
aggctcagggcgggctggctaacaattgctatattgaacaacaacttgaataccttgatccagcgttccaaccacac  
tcaggccgccaat (SEQ ID NO: 9) ;

15 DPA1\*020103 :

ccaatgltgcaacttatgccgcgtttgtacagacgcatagaccaacaggggagtttatgtttgaatttgatgaagat  
gagcAgttcttatgttgatctggacaagaaggagaccgtctggcatctggaggagtttggccgagccttttcccttg  
aggctcagggcgggctggctaacaattgctatattgaacaacaacttgaataccttgatccagcgttccaaccacac  
tcaggccgccaat (SEQ ID NO: 10) ;

20 DPA1\*020104 :

gcgtttgtacaaacccatagaccaacaggggagtttatgtttgaatttgatgaagatgagcagttcttatgttgatc  
tggataaAaaggagaccgtctggcatctggaggagtttggccgagccttttcccttgaggctcagggcgggctggc  
taacaattgctatattgaacaacaacttgaataccttgatccagcgttccaaccacac tcaggccgccaat (SEQ  
ID NO: 11) ;

25 DPA1\*020105 :

ccaatgltgcaacttatgccgcgtttgtacagacgcatagaccaacaggAgagtttatgtttgaatttgatgaagat  
gagcAgttcttatgttgatctggacaagaaggagaccgtctggcatctggaggagtttggccgagccttttcccttg



aggctcagggcgggctggctaacaatgctatattgaacaacaacttgaataccttgatccagcgltccaaccacac  
tcaggccgccaat(SEQ ID NO: 1 2) ;

DPA1\*020106 :

ccatgtgtcaacttatgccggtttgtacagacCcatagaccaacaggggagtttatgtttgaatttgatgaagat  
5 gagcagttctatgtggatctggataagaaggagaccgctcggcatctggaggagtttggccgagccttttcccttg  
aggctcagggcgggctggctaacaatgctatattgaacaacaacttgaataccttgatccagcgltccaaccacac  
tcaggccgccaat(SEQ ID NO: 1 3) ;

DPA1\*020201 :

aacttatgccatgtttgtacagacccatagaccaacaggAgagtttatgtttgaatttgatgaagatgagcagttc  
10 tatgtggatctggataagaaggagaccgctcggcatctggaggagtttggccgagccttttcccttgaggctcagg  
gctgggctggctaacaatgctatattgaacaacaacttgaataccttgatccagcgltccaaccacactcaggccgc  
caaT(SEQ ID NO: 1 4) ;

DPA1\*020202 :

ccatgtgtcaacttatgccatgtttgtacagacCcatagaccaacaggAgagtttatgtttgaatttgatgaagat  
15 gagcAgttctatgtggatctggataagaaggagaccgctcggcatctggaggagtttggccgagccttttcccttg  
aggctcagggcgggctggctaacaatgctatattgaacaacaacttgaataccttgatccagcgltccaaccacac  
tcaggccgccaat(SEQ ID NO: 1 5) ;

DPA1\*020203 :

atgtgtcaacttatgccaTgtttgtacagacccatagaccaacaggggagtttatgtttgaatttgatgaagaiga  
20 gcagttctatgtggatctggataagaaggagaccgctcggcatctggaggagtttggccgagccttttcccttgag  
gctcagggcgggctggctaacaatgctatattgaacaacaacttgaataccttgatccagcgltccaaccacactc  
aggccgccaat(SEQ ID NO: 1 6) ;

DPA1\*0203 :

ccatgtgtcaacttatgccggtttgtacagacCcatagaccaacaggggagtttatgtttgaatttgatgaagat  
25 gagatgttctatgtggatctggataagaaggagaccgctcggcatctggaggagtttggccgagccttttcccttg  
aggctcagggcgggctggctaacaatgctatattgaacaacaacttgaataccttgatccagcgltccaaccacac  
tcaggccgccaat(SEQ ID NO: 1 7) ;

DPA1\*0301 :

gccatgtttgtacagacccatagaccaacaggggagtttatgtttgaatttgaatgaagatgagatgttctatgttg  
atctggacaagaaggagaccgtctggcatctggaggagtttggccaagccttttccctttgaggctcagggcgggct  
ggctaacattgctatatCgaacaacaacttgaataccttgatccagcgttccaaccacactcaggccaccaac (SE

5 Q ID NO: 18) ;

DPA1\*0302 :

ccatgtgtcaacttatgccatgtttgtacagacccatagaccaacaggggagtttatgtttgaatttgaatgaagat  
gagatgttctatgttgatctggacaagaaggagaccgtctggcatctggaggagtttggccaagccttttccctttg  
aggctcagggcgggctggctaacattgctatatitgaacaacaacttgaataccttgatccagcgttccaaccacac

10 tcaggccaccaac (SEQ ID NO: 19) ;

DPA1\*0401 :

gccgcgtttgtacagacgcatalagaacaacaggagagtttatgtttgagtttgatgatgatgagatgttctatgttg  
atctggacaagaaggagaccgtctggcatctggaggagtttggccgagccttttccctttgaggctcagggcgggct  
ggctaacattgctatatitgaacaacaacttgaatatcgctatccagcgttccaaccacactcaggccgccaat (SE

15 Q ID NO: 20) ;

DPB1\*010101 :

agaattacgtgtaccaggacggcaggaatgctacgcgtttaatgggacacagcgttccctggagagatacatctta  
caaccgggaggagtagcgcgcttcgacagcgacgtgggggagttccgggcggtagcgagctggggcggcctgct  
gcggagtagtgaacagccagaaggacatcctggaggagaagcgggcagtgccggacaggGtAtgcagacacaact

20 acgagctggacgagggcgtgacctgcagcgccgagtcc (SEQ ID NO: 21) ;

DPB1\*010102 :

aatlacgtgtaccaggacggcaggaatgctacgcgtttaatgggacacagcgttccctggagagatacatctaca  
accgggaggagtagcgcgcttcgacagcgacgtgggAgagttccgggcggtagcgagctggggcggcctgctgc  
ggagtagtgaacagccagaaggacatcctggaggagaagcgggcagtgccggacaggGtAtgcagacacaactac

25 gtagctggacgagggcgtgacctgcagcgccga (SEQ ID NO: 22) ;

DPB1\*020102 (SEQ ID NO: 23) :

agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgttccctggagagatacatctta

caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gAggagttacttggaacagccagaaggacatcctggaggagGagcgggcagtgccggacaggatGtgcagacacaact  
acgagctggGcgggccccatgacctgcagcgccgagttcc (SEQ ID NO: 2 4) ;

DPB1\*020103 :

- 5 agaattaccttttccagggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgaC  
gaggagttacttggaacagccagaaggacatcctggaggaggagcgggcagtgccggacaggatgtgcagacacaact  
acgagctgggcgggccccatgacctgcagcgccgag (SEQ ID NO: 2 5) ;

DPB1\*020104 :

- 10 agaattaccttttccagggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggagttacttggaacagccagaaggacatcctggaggaggagcgggcagtgccggacaggatgtgcagacacaact  
acgagctgggcgggccccatgacctgcagcgccga (SEQ ID NO: 2 6) ;

DPB1\*020105 :

- 15 agaattaccttttccagggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaAgagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggagttacttggaacagccagaaggacatcctggaggaggagcgggcagtgccggacaggatgtgcagacacaact  
acgagctgggcgggccccatgacctgcagcgccgag (SEQ ID NO: 2 7) ;

DPB1\*020106 :

- 20 agaattaccttttccagggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaggagttTgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggagttacttggaacagccagaaggacatcctggaggaggagcgggcagtgccggacaggatgtgcagacacaact  
acgagctgggcgggccccatgacctgcagcgccgag (SEQ ID NO: 2 8) ;

DPB1\*0202 :

- 25 agaattaccttttccagggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaggagCtcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgaG  
gcggagttacttggaacagccagaaggacatcctggaggagGagcgggcagtgccggacaggatgtgcagacacaact

acgagctgggcgggcccAtgacctgcagcgccgag (SEQ ID NO: 29) ;

DPB1\*030101 :

agaattacgtgtaccagtTaccgcaggaatgctacgcgtttaatgggacacagcgcttccTggagagatacatcta  
caaccgggaggagttcgTgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccTgat  
5 gaggactacttggaacagccagaaggacCtccTggaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgagtc (SEQ ID NO: 30) ;

DPB1\*030102 :

agaattacgtgtaccagtTaccgcaggaatgctacgcgtttaatgggacacagcgcttccTggagagatacatcta  
caaccgggaggagttcgTgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccTgat  
10 gaggactacttggaacagccagaaggacctccTggaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctAcagcgccgag (SEQ ID NO: 31) ;

DPB1\*0401 :

agaattaccittttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccTggagagatacatcta  
caaccgggaggagttcgTgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccTgct  
15 gcggagtacttggaacagccagaaggacatccTggaggagaagcgggcagtgccggacaggatGtcagacacaact  
acgagctggGcgggcccattgacctgcagcgccgagtc (SEQ ID NO: 32) ;

DPB1\*0402 :

agaattaccittttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccTggagagatacatcta  
caaccgggaggagttcgTgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccTgat  
20 gAggagtacttggaacagccagaaggacatccTggaggagaagcgggcagtgccggacaggatGtcagacacaact  
acgagctggGcgggcccattgacctgcagcgccgagtc (SEQ ID NO: 33) ;

DPB1\*0501 :

agaattaccittttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccTggagagatacatcta  
caaccgggaggagCtcgTgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccTgaG  
25 gcggagtacttggaacagccagaaggacatccTggaggagaagcgggcagtgccggacaggatGtcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 34) ;

DPB1\*0601 :

agaattacgtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccitggagagatacatctta  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgat  
gaggaCtactggaacagccagaaggacCtcttggaggagGagcgggcagtgccggacaggatGtgcagacacaact  
acgagctggacgaggccgtgacctgcag (SEQ ID NO: 3 5) ;

5 DPB1\*0801 :

cttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccitggagagatacatcttaacaaccggg  
aggagttcgtTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgatgAggagta  
ctggaacagccagaaggacatccttggaggagGagcgggcagtgccggacagggtatgcagacacaacttacgagctg  
gacgaggccgtgacctgcag (SEQ ID NO: 3 6) ;

10 DPB1\*0901 :

agaattacgtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccitggagagatacatctta  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgat  
gaggaCtactggaacagccagaaggacatccttggaggagGagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 3 7) ;

15 DPB1\*1001 :

agaattacgtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccitggagagatacatctta  
caaccgggaggagttcgtTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgat  
gAggagttactggaacagccagaaggacatccttggaggagGagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 3 8) ;

20 DPB1\*110101 :

gtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccitggagagatacatcttaacaaccggC  
aggagttacgcgcgttcgacagcgacgtgggagagttccgggcggtgacggagctggggcgccctgctgcggagta  
ctggaacagccagaaggacctccttggaggagaggcgggcagtgccggacaggatgtgcagacacaacttacgagctg  
gacgaggccgtgacctgcag (SEQ ID NO: 3 9) ;

25 DPB1\*110102 :

agaattacgtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccitggagagatacatctta  
caacAggcaggagttacgcgcgttcgacagcgacgtgggagagttccgggcggtgacggagctggggcgccctgct

gcggagtagtggaaacagccagaaggacctccaggaggagaggcgggcagtgccggacaggatgtgcagacacaact  
acgagctggacgaggccgtgacctgcag (SEQ ID NO: 4 0) ;

DPB1\*1301 :

agaattacgtgtaccagtTaccggcaggaatgctacgcgtttaatgggacacagcgcttccaggagatacatctta  
5 caaccgggaggagtagcgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgct  
gcggagtagtggaaacagccagaaggacatccaggaggagGagcgggcagtgccggacaggAtatgcagacacaact  
acgagctggacgaggccgtgacctgcag (SEQ ID NO: 4 1) ;

DPB1\*1401 :

agaattacgtgcaccagtTaccggcaggaatgctacgcgtttaatgggacacagcgcttccaggagatacatctta  
10 caaccgggaggagttcgTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggaCtagtggaaacagccagaaggacCtccaggaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcag (SEQ ID NO: 4 2) ;

DPB1\*1501 :

agaattacgtgtaccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccaggagatacatctta  
15 caaccggCaggagtagcgcgcttcgacagcgacgtgggagagttccgggcggtgacggagctggggcggcctgct  
gcggagtagtggaaacagccagaaggacctccaggaggagaggcgggcagtgccggacaggatgtgcagacacaact  
acgagctgggtcgggcccAtgacctgcagcgccgag (SEQ ID NO: 4 3) ;

DPB1\*1601 :

agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccaggagatacatctta  
20 caaccgggaggagttcgTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gAggagtagtggaaacagccagaaggacatccaggaggagGagcgggcagtgccggacaggatGtgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 4 4) ;

DPB1\*1701 :

agaattacgtgcaccagtTaccggcaggaatgctacgcgtttaatgggacacagcgcttccaggagatacatctta  
25 caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggaCtagtggaaacagccagaaggacatccaggaggagGagcgggcagtgccggacaggatGtgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 4 5) ;

DPB1\*1801 :

gtgtaccagggacggcaggaatgctacgcgtttaatgggacacagcgcttccitggagagatacatctacaaccggg  
aggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtagcgagctggggcggcctgatgAggagta  
ctggaacagccagaaggacatccitggaggagaagcgggcagtgccggacaggatgtgcagacacaactacgagctg  
5 gTcgggcccattgacctgcag(SEQ ID NO: 4 6) ;

DPB1\*1901 :

agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccitggagagatacatctta  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtagcgagctggggcggcctgaG  
gcggagttactggaacagccagaaggacatccitggaggagGagcgggcagtgccggacaggAtatgcagacacaact  
10 acgagctggacgaggccgtgacctgcagcgccgag(SEQ ID NO: 4 7) ;

DPB1\*200101 :

agaattacgtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccitggagagatacatctta  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtagcgagctggggcggcctgat  
gaggactactggaacagccagaaggacCtccitggaggagaagcgggcagtgccggacaggatGtgcagacacaact  
15 acgagctggacgaggccgtgacctgcagcgccgag(SEQ ID NO: 4 8) ;

DPB1\*200102 :

agaattacgtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccitggagagatacatctta  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtagcgagctggggcggcctgat  
gaggactactggaacagccagaaggacctccitggaggagaagcgggcagtgccggacaggatgtgcagacacaact  
20 acgagctggacgaggccgtgacctgcagcgTcga(SEQ ID NO: 4 9) ;

DPB1\*2101 :

agaattacgtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccitggagagatacatctta  
caaccgggaggagCtgcgttcgacagcgacgtgggggagttccgggcggtagcgagctggggcggcctgaG  
gcggagttactggaacagccagaaggacatccitggaggagGagcgggcagtgccggacaggatGtgcagacacaact  
25 acgagctggacgaggccgtgacctgcagcgccgag(SEQ ID NO: 5 0) ;

DPB1\*2201 :

agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccitggagagatacatctta

caaccgggaggagCtcgtgcgcttcgacagcgacgtgggggagtccgggcggtgacggagctggggcgccctgaG  
gcggagtlactggaacagccagaaggacatccctggaggagGagcgggcagtgccggacaggatGtcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 5 1) ;

DPB1\*2301 :

- 5 agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttcctggagagatacatctta  
caaccgggaggagttcgTgcgcttcgacagcgacgtgggggagtccgggcggtgacggagctggggcgccctgct  
gcggagtlactggaacagccagaaggacatccctggaggagaagcgggcagtgccggacaggatGtcagacacaact  
acgagctggGcgggcccattgacctgcagcgccgag (SEQ ID NO: 5 2) ;

DPB1\*2401 :

- 10 agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttcctggagagatacatctta  
caaccgggaggagttcgCgcgcttcgacagcgacgtgggggagtccgggcggtgacggagctggggcgccctgaG  
gcggagtlactggaacagccagaaggacatccctggaggagaagcgggcagtgccggacaggatGtcagacacaact  
acgagctgggCgggcccAtgacctgcagcgccgag (SEQ ID NO: 5 3) ;

DPB1\*2501 :

- 15 agaattacgtgtaccagTtacggcaggaatgctacgcgtttaatgggacacagcgcttcctggagagatacatctta  
caaccgggaggagttcgTgcgcttcgacagcgacgtgggggagtccgggcggtgacggagctggggcgccctgat  
gAggagtlactggaacagccagaaggacCtccctggaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 5 4) ;

DPB1\*260101 :

- 20 ggtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttcctggagagatacatcttaaccggg  
aggagtlacgcgcgcttcgacagcgacgtgggagagtccgggcggtgacggagctggggcgccctgctgcggagta  
ctggaacagccagaaggacatccctggaggagaagcgggcagtgccggacagAgtaTgcagacacaactacgagctg  
gacgaggccgtgacctgcagcgccgag (SEQ ID NO: 5 5) ;

DPB1\*260102 :

- 25 ggtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttcctggagagatacatcttaaccggg  
aggagtlacgcgcgcttcgacagcgacgtgggggagtccgggcggtgacggagctggggcgccctgctgcggagta  
ctggaacagccagaaggacatccctggaggagaagcgggcagtgccggacagggtatgcagacacaactacgagctg



gacgaggccgtgacctgcagcgccga (SEQ ID NO: 56) ;

DPB1\*2701 :

agaattacgtgtaccagTTacggcaggaatgctacgcgtttaatgggacacagcgcttccTggagagatacatcta  
caaccgggaggagTtcgcgcgttcgacagcgacgtgggggagTtccgggcggTgacggagctggggcgccTgct  
5 gcgagTacttggaacagccagaaggacatccTggaggagaagcgggcagTgccggacaggatGtgacacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 57) ;

DPB1\*2801 :

agaattacctttccagggacggcaggaatgctacgcgtttaatgggacacagcgcttccTggagagatacatcta  
caaccgggaggagTtcgcgcgttcgacagcgacgtgggggagTtccgggcggTgacggagctggggcgccTgat  
10 gAggagTacttggaacagccagaaggacCtccTggaggagaagcgggcagTgccggacaggatGtgacacacaact  
acgagctggTcgggcccatgacctgcagcgccgag (SEQ ID NO: 58) ;

DPB1\*2901 :

agaattacgtgtaccagTtacggcaggaatgctacgcgtttaatgggacacagcgcttccTggagagatacatcta  
caaccgggaggagTtcgtgcgttcgacagcgacgtgggggagTtccgggcggTgacggagctggggcgccTgat  
15 gaggaCtacttggaacagccagaaggacCtccTggaggagGagcgggcagTgccggacaggatGtgacacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 59) ;

DPB1\*3001 :

agaattacgtgcaccagTTacggcaggaatgctacgcgtttaatgggacacagcgcttccTggagagatacatcta  
caaccgggaggagTtcgtgcgttcgacagcgacgtgggggagTtccgggcggTgacggagctggggcgccTgaG  
20 gcgagTacttggaacagccagaaggacatccTggaggagGagcgggcagTgccggacaggatGtgacacacaact  
acgagctggacgaggccgtgacctgcag (SEQ ID NO: 60) ;

DPB1\*3101 :

agaattacctttccagggacggcaggaatgctacgcgtttaatgggacacagcgcttccTggagagatacatcta  
caaccgggaggagTtcgcgcgttcgacagcgacgtgggggagTtccgggcggTgacggagctggggcgccTgct  
25 gcgagTacttggaacagccagaaggacctccTggaggagaagcgggcaTtgccggacaggatGtgacacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 61) ;

DPB1\*3201 :

agaattaccittttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggTgtacttggaacagccagaaggacatccctggaggaggagcgggcagtgccggacaggatgtgcagacacaact  
acgagctgggcgggcccattgacctgcagcgccgag (SEQ ID NO: 6 2) ;

5 DPB1\*3301 :

agaattaccittttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaggagttcgcgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgct  
gcggagtacttggaacagccagaaggacatccctggaggagGagcgggcagtgccggacaggatGtgcagacacaact  
acgagctggGcgggcccattgacctgcag (SEQ ID NO: 6 3) ;

10 DPB1\*3401 :

agaattaccittttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaggagctcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgct  
gcggagtacttggaacagccagaaggacctccctggaggagaagcgggcattgcccggacaggatgtgcagacacaact  
acgagctggtcgggcccAttgacctgcag (SEQ ID NO: 6 4) ;

15 DPB1\*3501 :

agaattacgtgtaccagtTaccgcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaggagttcgTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggaCtacttggaacagccagaaggacatccctggaggagaagcgggcagtgccggacaggatgtgcagacacaact  
acgagctggacgaggccgtgacctgcag (SEQ ID NO: 6 5) ;

20 DPB1\*3601 :

agaattacgtgtaccagtTaccgcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaggagCtcgTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgaG  
gcggagtacttggaacagccagaaggacatccctggaggagaagcgggcagtgccggacaggatGtgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 6 6) ;

25 DPB1\*3701 :

gtgtaccagttaccgcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctacaaccggg  
aggagttcgTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgatgAggagta

ctggaacagccagaaggacatccctggaggagGagcgggcagtgccggacagggtatgcagacacaactacgagctg  
gacgaggccgtgacctgcagcgccgag (SEQ ID NO: 67) ;

DPB1\*3801 :

cttttccaggacggcaggaatgctacCcgittaatgggacacagcgcttccctggagagatacatctacaaccggg  
5 aggagctcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgaggcggagta  
ctggaacagccagaaggacatccctggaggagaagcgggcagtgccggacaggatgtgcagacacaactacgagctg  
gacgaggccgtgacctgcag (SEQ ID NO: 68) ;

DPB1\*3901 :

agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctla  
10 caaccgggaggagtagcgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgct  
gcggagtagtggaaacagccagaaggacatccctggaggagaagcgggcagtgccggacaggatGtgcagacacaact  
acgagctggGcgggcccatgacctgcagcgccga (SEQ ID NO: 69) ;

DPB1\*4001 :

agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctla  
15 caaccgggaggagtagcgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgct  
gcggagtagtggaaacagccagaaggacatccctggaggagaagcgggcagtgccggacaggatgtgcagacacaact  
acgagctggTcgggcccatgacctgcagcgccga (SEQ ID NO: 70) ;

DPB1\*4101 :

aattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctla  
20 accgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgatlga  
ggagtagtggaaacagccagaaggacTtccctggaggagGagcgggcagtgccggacaggatgtgcagacacaactac  
gagctgggcgggcccatgacctgcagcgccga (SEQ ID NO: 71) ;

DPB1\*4401 :

agaattacgtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctla  
25 caaccgggaggagCtcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgatlga  
gaggacttagtggaaacagccagaaggacCtccctggaggagGagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 72) ;

DPB1\*4501 :

gtgcaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctacaaccggg  
aggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtagcggagctggggcggcctgatgAggagta  
ctggaacagccagaaggacCtccctggaggagaagcgggcagtgccggacagggtatgcagacacaactacgagctg  
5 gacgaggccgtgacctgcag (SEQ ID NO: 7 3) ;

DPB1\*4601 :

agaattaccctttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatct  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtagcggagctggggcggcctgat  
gaggaCtactggaacagccagaaggacatccctggaggagGagcgggcagtgccggacaggatgtgcagacacaact  
10 acgagctgggcgggcccAtgacctgcagcgccgag (SEQ ID NO: 7 4) ;

DPB1\*4701 :

agaattaccctttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatct  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtagcggagctggggcggcctgaG  
gcggagtactggaacagccagaaggacatccctggaggagGagcgggcagtgccggacaggatgtgcagacacaact  
15 acgagctgggcgggcccAtgacctgcagcgccgag (SEQ ID NO: 7 5) ;

DPB1\*4801 :

aattaccctttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctaca  
accgggaggagCtcgtgcgttcgacagcgacgtgggggagttccgggcggtagcggagctggggcggcctgatgA  
ggagtlactggaacagccagaaggacatccctggaggaggagcgggcagtgccggacaggatgtgcagacacaactac  
20 gagctggGcgggcccAtgacctgcag (SEQ ID NO: 7 6) ;

DPB1\*4901 :

aattaccctttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctaca  
accgggaggagtagcgcgttcgacagcgacgtgggggagttccgggcggtagcggagctggggcggcctgatgA  
ggagtlactggaacagccagaaggacatccctggaggagaagcgggcagtgccggacaggatGtgcagacacaactac  
25 gagctggGcgggcccAtgacctgcag (SEQ ID NO: 7 7) ;

DPB1\*5001 :

aattacgtgtaccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctaca

accgggaggagttcgTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgaatga  
ggaCtacttggaacagccagaaggacCtcttgaggagaagcgggcagtgccggacagggtatgcagacacaactac  
gagctggacgaggccgtgacctgcag (SEQ ID NO: 78) ;

DPB1\*5101 :

- 5 agaattaccittttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgaat  
gAggagttacttggaacagccagaaggacatcttgaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggGcgggcccatgacctgcagcgccgag (SEQ ID NO: 79) ;

DPB1\*5201 :

- 10 agaattacgtgtaccagtTaccggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgct  
gcggagtacttggaacagccagaaggacCtcttgaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcag (SEQ ID NO: 80) ;

DPB1\*5301 :

- 15 agaattaccittttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttacgcgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgaat  
gAggagttacttggaacagccagaaggacatcttgaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggTcgggcccatgacctgcag (SEQ ID NO: 81) ;

DPB1\*5401 :

- 20 agaattacgtgcaccagtTaccggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgaG  
gcggagtacttggaacagccagaaggacatcttgaggagGagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcag (SEQ ID NO: 82) ;

DPB1\*5501 :

- 25 agaattacgtgcaccagtTaccggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgct  
gcggagtacttggaacagccagaaggacatcttgaggagGagcgggcagtgccggacagggtatgcagacacaact

acgagctggacgaggccgtgacctgcag (SEQ ID NO: 8 3) ;

DPB1\*5601 :

gtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctacaaccggg  
aggagtTcgcgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgctgcggagta  
5 ctggaacagccagaaggacCtccctggaggagaagcgggcagtgccggacagggtatgcagacacaactacgagctg  
gacgaggccgtgacctgcag (SEQ ID NO: 8 4) ;

DPB1\*5701 :

cttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctacaaccggg  
aggagttcgTcgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgatgaggaCta  
10 ctggaacagccagaaggacCtccctggaggagaagcgggcagtgccggacagggtatgcagacacaactacgagctg  
gacgaggccg (SEQ ID NO: 8 5) ;

DPB1\*5801 :

aattacgtgcaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctaca  
accgggaggagCtcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgctgc  
15 ggagttactggaacagccagaaggacatccctggaggagGagcgggcagtgccggacaggatGtgcagacacaactac  
gagctggacgaggccgtgacctgcag (SEQ ID NO: 8 6) ;

DPB1\*5901 :

agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctaca  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgat  
20 gAggagttactggaacagccagaaggacCtccctggaggagaagcgggcagtgccggacaggatGtgcagacacaact  
acgagctggGcgggcccattgacctgcag (SEQ ID NO: 8 7) ;

DPB1\*6001 :

agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctaca  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgat  
25 gaggagttactggaacagccagaaggacaAccctggaggagaagcgggcagtgccggacaggatGtgcagacacaact  
acgagctggGcgggcccattgacctgcag (SEQ ID NO: 8 8) ;

DPB1\*6101N :

agaattacgtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggactactggaacagccagaaggacctcttTaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgc (SEQ ID NO: 89) ;

5 DPB1\*6201 :

agaattacctttccagggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagCtcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgct  
gcggagtactggaacagccagaaggacatcttggaggagaagcgggcagtgccggacaggatGtcagacacaact  
acgagctggTcgggcccattgacctgcag (SEQ ID NO: 90) ;

10 DPB1\*6301 :

aattacctttccagggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctaca  
accgggaggagCtcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgctgc  
ggagtactggaacagccagaaggacatcttggaggagaagcgggcagtgccggacaggatGtcagacacaactac  
gagctggacgaggccgtgacctgcag (SEQ ID NO: 91) ;

15 DPB1\*6401N :

aattaaagtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctaca  
accgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgatga  
ggaCtactggaacagccagaaggacCtccctggaggagGagcgggcagtgccggacaggatGtcagacacaactac  
gagctggacgaggccgtgacctgcag (SEQ ID NO: 92) ;

20 DPB1\*6501 :

agaattacctttccagggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttacgcgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgct  
gcggagtactggaacagccagaaggacatcttggaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgcgag (SEQ ID NO: 93) ;

25 DPB1\*6601 :

agaattacgtgcaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgcgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgct

gcggagtagctggaacagccagaaggacatcctggaggagaagcgggcagtgccggacaggatGtcagacacaact  
acgagctggGcggggcccatgacctgcagcgccgag (SEQ ID NO: 9 4) ;

DPB1\*6701 :

agaattacgtgcaccagTaccggcaggaatgctacgcgtttaatgggacacagcgcttcctggagagatacatcta  
5 caaccgggaggagtagTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgct  
gcggagtagctggaacagccagaaggacCtctggaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 9 5) ;

DPB1\*6801 :

agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttcctggagagatacatcta  
10 caaccgggaggagtagTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgat  
gAggagtagctggaacagccagaaggacatcctggaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccga (SEQ ID NO: 9 6) ;

DPB1\*6901 :

agaattacgtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttcctggagagatacatcta  
15 caaccgggaggagtagTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgat  
gaggaCtagctggaacagccagaaggacctcctggaggagaGcggggcagtgccggacaggatgtgcagacacaact  
acgagctggacgaggccgtgacc (SEQ ID NO: 9 7) ;

DPB1\*7001 :

aattacgtggaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttcctggagagatacatctaca  
20 accgggaggagtagTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgatga  
ggaCtagctggaacagccagaaggacCtctggaggagaagcgggcagtgccggacagggtatgcagacacaactac  
gagctggacgaggccgtgacctgcag (SEQ ID NO: 9 8) ;

DPB1\*7101 :

aattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttcctggagagatacatctaca  
25 accgggaggagtagTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgctgc  
ggagtagctggaacagccagaaggacatcctggaggagGagcgggcagtgccggacaggatGtcagacacaactac  
gagctggGcggggcccatgacctgcag (SEQ ID NO: 9 9) ;



DPBI\*7201 :

aattaccctttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctaca  
accgggaggagttcgcgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgctgc  
ggagtactggaacagccagaaggacCtctggaggagaagcgggcagtgccggacaggatGtcagacacaactac  
5 gagctggGcgggccccatgaccctgcag(SEQ ID NO: 1 0 0) ;

DPBI\*7301 :

aattaccctttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctaca  
accgggaggagttcgctgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgatGA  
ggagtactggaacagccagaaggacCtctggaggagaagcgggcagtgccggacaggatGtcagacacaactac  
10 gagctggGcgggccccatgaccctgcag(SEQ ID NO: 1 0 1) ;

DPBI\*7401 :

gtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctacaaccggC  
aggagtacgcgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgctgcggagta  
ctggaacagccagaaggacctcctggaggagaggcgggcagtgccggacaggatgtgcagacacaactacgagctg  
15 gtccggccccAtgaccctgcag(SEQ ID NO: 1 0 2) ;

DPBI\*7501 :

cttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctacaaccggg  
aggagttcgTgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgatGaggagta  
ctggaacagccagaaggacatcctggaggagaagcgggcagtgccggacagggtatgcagacacaactacgagctg  
20 gGcgggccccatgaccctgcag(SEQ ID NO: 1 0 3) ;

DPBI\*7601 :

agaattacgtgcaccagtTaccggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctac  
caaccgggaggagttcgcgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggaCtactggaacagccagaaggacCtctggaggagaagcgggcagtgccggacagggtatgcagacacaact  
25 acgagctggacgaggccgtgaccctgcag(SEQ ID NO: 1 0 4) ;

DPBI\*7701 :

agaattaccctttccaggacTgcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctac

caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggagttactggaacagccagaaggacatcctggaggagaagcgggcagtgccggacaggatgtgcagacacaact  
acgagctggggcgggcccattgacctgcagcgccgag (SEQ ID NO: 105) ;

DPB1\*7801 :

- 5 agaattacgtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggactactggaacagccagaaggacctcctggaggagaagcgggcagtgctggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 106) ;

DPB1\*7901 :

- 10 agaattacgtgtaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggagttactggaacagccagaaggacatcctggaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 107) ;

DPB1\*8001 :

- 15 agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggactactggaacagccagaaggacatcctggaggagaagcgggcagtgccggacaggatgtgcagacacaact  
acgagctgggcgggcccattgacc (SEQ ID NO: 108) ;

DPB1\*8101 :

- 20 agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggagttactggaacagccagaaggacatcctggaggagGagcgggcagtgccggacaggatgtgcagacacaact  
acgagctggGcgggcccattgacctgcagcgccgag (SEQ ID NO: 109) ;

DPB1\*8201 :

- 25 agaattaccttttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggagttactggaacagccagaaggacatcctggaggagaagcgggcagtgccggacaggatgtgcagacacaact

acgagctgggcgggcccAtgacctgcagcAccgag(SEQ ID NO: 1 1 0) ;

DPBI\*8301 :

agaattacctttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgat  
5 gaggagtactggaacagccagaaggacTtccctggaggagaagcgggcagtgccggacaggatgtgcagacacaact  
acgagctgggcgggccccatgacctgcagcgccgag(SEQ ID NO: 1 1 1) ;

DPBI\*8401 :

agaattacctttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgaG  
10 gAggagtactggaacagccagaaggacatccctggaggagaagcgggcagtgccggacaggatgtgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccga(SEQ ID NO: 1 1 2) ;

DPBI\*8501 :

agaattacgtgtaccagtTaccggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgct  
15 gcgaggtactggaacagccagaaggacatccctggaggagaagcgggcagtgccggacaggatgtgcagacacaact  
acgagctggacgaggccgtgacctgcagcAccgag(SEQ ID NO: 1 1 3) ;

DPBI\*8601 :

gaattacgtgcaccagtTaccggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctac  
aaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgatg  
20 aggaCtactggaacagccagaaggacatccctggaggagGagcgggcagtgccggacaggatgtgcagacacaacta  
cgagctgggcgggcccAtgacctgcagcgccga(SEQ ID NO: 1 1 4) ;

DPBI\*8701 :

agaattacgtgtaccagtTaccggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatcta  
caaccgggaggagttcgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgct  
25 gcgaggtactggaacagccagaaggacCtccctggaggagaagcgggcagtgccggacaggatGtgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag(SEQ ID NO: 1 1 5) ;

DPBI\*8801 :

agaattacgtgtaccagtTaccggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggaCtactggaacagccagaaggacatccctggaggagGagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag(SEQ ID NO: 1 1 6) ;

5 DPBI\*8901 :

agaattacgtgtaccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgct  
gcggagttactggaacagccagaaggacatccctggaggagaagcgggcagtgccggacaggatGtcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag(SEQ ID NO: 1 1 7) ;

10 DPBI\*9001 :

agaattacgtgtaccaggacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgct  
gcggagttactggaacagccagaaggacatccctggaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag(SEQ ID NO: 1 1 8) ;

15 DPBI\*9101 :

agaattacgtgcaccagttacggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggaCtactggaacagccagaaggacCtccctggaggagaagcgggcagtgccggacagggtatGtcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag(SEQ ID NO: 1 1 9) ;

20 DPBI\*9201 :

agaattacgtgtaccagtTaccggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat  
gaggaCtactggaacagccagaaggacCtccctggaggagaagcgggcagtgccggacagggtatgcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag(SEQ ID NO: 1 2 0) ;

25 DPBI\*9301 :

agaattacgtgtaccagtTaccggcaggaatgctacgcgtttaatgggacacagcgcttccctggagagatacatctta  
caaccgggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgat

gAggagtagtggaacagccagaaggacatccaggaggagGagcgggcagtgccggacaggatGtcagacacaact  
acgagctggacgaggccgtgacctgcagcgccgag (SEQ ID NO: 1 2 1) ;

DPB1\*9601 :

agaattacctttccaggacggcaggaatgctacgcgtttaatgggacacagcgcttcctggagagatacatcta  
5 caaccgggaggagtagcgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccctgct  
gcggagtagtggaacagccagaagCacatccaggaggagaagcgggcagtgccggacaggatgtgcagacacaact  
acgagctgggcgggcccattgacctgcagcgccgag (SEQ ID NO: 1 2 2) ;

In the following, Probe lists DP1-DP4 are shown  
10 in Tables 13-1 to 13-3 and Tables 14-1 to 14-3  
respectively. Probe-Allele Lists DP1-4 are shown in  
Tables 15-1 to 15-5 and Tables 16-1 to 16-5.

Table 13-1

Probe No.	Base Sequence
0	acg cat aga cca aca ggG ( SEQ ID No: 123)
1	ag ttt atg ttt gaa ttt gat gaA ( SEQ ID No: 124)
2	t ctg gag gag ttt ggc cA ( SEQ ID No: 125)
3	g acg cat aga cca aca ggA ( SEQ ID No: 126)
4	g ttt atg ttt gaa ttt gat gaC ( SEQ ID No: 127)
5	cac act cag gcc gcc aaT ( SEQ ID No: 128)
6	ttc tat gtg gat ctg gat aaA ( SEQ ID No: 129)
7	ctg gag gag ttt ggc caa A ( SEQ ID No: 130)
8	ctg gag gag ttt ggc cG ( SEQ ID No: 131)
9	gcc gcg ttt gta cag acC ( SEQ ID No: 132)
10	t gaa ttt gat gaa gat gag cA ( SEQ ID No: 133)
11	ag ttc tat gtg gat ctg gaT ( SEQ ID No: 134)
12	g acc cat aga cca aca ggA ( SEQ ID No: 135)
13	t gcc atg ttt gta cag acC ( SEQ ID No: 136)
14	at gtg tca act tat gcc aT ( SEQ ID No: 137)
15	ctg gct aac att gct ata tC ( SEQ ID No: 138)
16	cat gtg tca act tat gcc aT ( SEQ ID No: 139)
17	aac aac aac ttg aat atc gcT ( SEQ ID No: 140)

Table 13-2

Probe No.	Base Sequence
0	gca gtg ccg gac agg G (SEQ ID No: 141)
1	ca gtg ccg gac agg gtA (SEQ ID No: 142)
2	tc gac agc gac gtg ggA (SEQ ID No: 143)
3	c aac cgg gag gag ttc gT (SEQ ID No: 144)
4	ctg ggg cgg cct gat gA (SEQ ID No: 145)
5	g gac atc ctg gag gag G (SEQ ID No: 146)
6	ca gtg ccg gac agg atG (SEQ ID No: 147)
7	a cac aac tac gag ctg gG (SEQ ID No: 148)
8	g ctg ggg cgg cct gaC (SEQ ID No: 149)
9	ag gag gag cgg gca gtT (SEQ ID No: 150)
10	ga tac atc tac aac cgg gaA (SEQ ID No: 151)
11	c tac aac cgg gag gag ttT (SEQ ID No: 152)
12	c tac aac cgg gag gag C (SEQ ID No: 153)
13	g ctg ggg cgg cct gaG (SEQ ID No: 154)
14	gag ctg ggc ggg ccc A (SEQ ID No: 155)
15	ag aat tac gtg tac cag tT (SEQ ID No: 156)
16	gg cgg cct gat gag gaC (SEQ ID No: 157)
17	gg aac agc cag aag gac C (SEQ ID No: 158)
18	ac gag gcc gtg acc ctA (SEQ ID No: 159)
19	c tac aac cgg gag gag tT (SEQ ID No: 160)
20	aac cgg gag gag ctc gT (SEQ ID No: 161)
21	g gac ctc ctg gag gag G (SEQ ID No: 162)
22	ag aat tac gtg cac cag tT (SEQ ID No: 163)
23	aga tac atc tac aac cgg C (SEQ ID No: 164)
24	g gag aga tac atc tac aac A (SEQ ID No: 165)
25	g gca gtg ccg gac agg A (SEQ ID No: 166)
26	gag ctg gtc ggg ccc A (SEQ ID No: 167)
27	ga cac aac tac gag ctg gT (SEQ ID No: 168)
28	cc gtg acc ctg cag cgT (SEQ ID No: 169)
29	gg gca gtg ccg gac agA (SEQ ID No: 170)
30	g gag gag aag cgg gca T (SEQ ID No: 171)

Table 13-3

Probe No.	Base Sequence
31	ggg cgg cct gat gag gT (SEQ ID No: 172)
32	ga cgg cag gaa tgc tac C (SEQ ID No: 173)
33	gg aac agc cag aag gac T (SEQ ID No: 174)
34	g gac ttc ctg gag gag G (SEQ ID No: 175)
35	gg aac agc cag aag gac aA (SEQ ID No: 176)
36	gc cag aag gac ctc ctg T (SEQ ID No: 177)
37	gac ctc ctg gag gag aG (SEQ ID No: 178)
38	aat tac ctt ttc cag gga cT (SEQ ID No: 179)
39	gag aag cgg gca gtg cT (SEQ ID No: 180)
40	ccc atg acc ctg cag cA (SEQ ID No: 181)
41	tg ggg cgg cct gag gA (SEQ ID No: 182)
42	gcc gtg acc ctg cag cA (SEQ ID No: 183)
43	g aat tac gtg cac cag tT (SEQ ID No: 184)
44	ac tgg aac agc cag aag C (SEQ ID No: 185)



Table 14-1

Probe No.	Base Sequence
0	a cca aca ggG gag ttt atg ( SEQ ID No : 186)
1	gaa ttt gat gaA gat gag atg ( SEQ ID No : 187)
2	ag ttt ggc cAa gcc ttt tc ( SEQ ID No : 188)
3	ga cca aca ggA gag ttt atg ( SEQ ID No : 189)
4	gaa ttt gat gaC gat gag atg ( SEQ ID No : 190)
5	at ctg gat aaA aag gag acc ( SEQ ID No : 191)
6	ttt ggc caa Acc ttt tcc tt (i SEQ ID No : 192)
7	ag ttt ggc cGa gcc ttt tc (i SEQ ID No : 193)
8	t gta cag acC cat aga cca ( SEQ ID No : 194)
9	gaa gat gag cAg ttc tat gt ( SEQ ID No : 195)
10	cg ttt gta caA acc cat aga ( SEQ ID No : 196)
11	g gat ctg gaT aag aag gag ( SEQ ID No : 197)
12	act tat gcc aTg ttt gta cag ( SEQ ID No : 198)
13	att gct ata tCg aac aac aac ( SEQ ID No : 199)
14	g aat atc gcT atc cag cgt ( SEQ ID No : 200)

Table 14-2

Probe No.

Base Sequence

0	tAc cag gga cgg cag ga (SEQ ID No: 201)
1	ccg gac agg Gta tgc aga (SEQ ID No: 202)
2	g gac agg gta tgc aga ca (SEQ ID No: 203)
3	gac gtg gga gag ttc cg (SEQ ID No: 204)
4	at tac ctt tTc cag gga cg (SEQ ID No: 205)
5	g gag ttc gTg cgc ttc g (SEQ ID No: 206)
6	gg cct gat gAg gag tac t (SEQ ID No: 207)
7	g gag gag GAg cgg gca (SEQ ID No: 208)
8	g gac agg atG tgc aga ca (SEQ ID No: 209)
9	gag ctg gGc ggg ccc (SEQ ID No: 210)
10	cgg cct gaC gag gag ta (SEQ ID No: 211)
11	cgg gca gtT ccg gac ag (SEQ ID No: 212)
12	c aac cgg gaA gag ttc gt (SEQ ID No: 213)
13	g gag gag ttT gtg cgc tt (SEQ ID No: 214)
14	g gag gag Ctc gtg cgc (SEQ ID No: 215)
15	cgg cct gaG gcg gag t (SEQ ID No: 216)
16	c ggg ccc Atg acc ctg (SEQ ID No: 217)
17	tg tac cag tTa cgg cag g (SEQ ID No: 218)
18	t gat gag gaC tac tgg aac (SEQ ID No: 219)
19	cag aag gac Ctc ctg gag (SEQ ID No: 220)
20	gtg acc ctA cag cgc cg (SEQ ID No: 221)
21	g gag gag tTc gcg cgc (SEQ ID No: 222)
22	g gag ctc gTg cgc ttc g (SEQ ID No: 223)
23	aat tac gtg Cac cag tta cg (SEQ ID No: 224)
24	tac aac cgg Cag gag tac (SEQ ID No: 225)
25	atc tac aac Agg cag gag t (SEQ ID No: 226)
26	ccg gac agg Ata tgc aga (SEQ ID No: 227)
27	c gag ctg gTc ggg ccc (SEQ ID No: 228)
28	g ccg gac agA gta tgc ag (SEQ ID No: 229)
29	g cac cag tTa cgg cag g (SEQ ID No: 230)
30	g cgg gca Ttg ccg gac (SEQ ID No: 231)

Table 14-3

Probe No.	Base Sequence
31	ct gat gag gTg tac tgg aa (SEQ ID No : 2 3 2)
32	gaa tgc tac Ccg ttt aat gg (SEQ ID No : 2 3 3)
33	cag aag gac Ttc ctg gag (SEQ ID No : 2 3 4)
34	ag aag gac aAc ctg gag g (SEQ ID No : 2 3 5)
35	gac ctc ctg Tag gag aag (SEQ ID No : 2 3 6)
36	g gag gag aGg cgg gca (SEQ ID No : 2 3 7)
37	g gac cag tTa cgg cag g (SEQ ID No : 2 3 8)
38	tc cag gga cTg cag gaa t (SEQ ID No : 2 3 9)
39	g gca gtg cTg gac agg g (SEQ ID No : 2 4 0)
40	g ctg ggc gGg ccc atg (SEQ ID No : 2 4 1)
41	cgg cct gaG gag gag ta (SEQ ID No : 2 4 2)
42	gg cct gag gAg gag tac t (SEQ ID No : 2 4 3)
43	agc cag aag Cac atc ctg (SEQ ID No : 2 4 4)

Table 15-1

Allele Number		Probe Number for Detection			
DPA1*010301	0	1	2		
DPA1*010302	3				
DPA1*0104	4				
DPA1*0105	5				
DPA1*0106	6				
DPA1*0107	7				
DPA1*0108	4	8			
DPA1*020101	9	6	5		
DPA1*020102	6	5			
DPA1*020103	10	5			
DPA1*020104	6	5			
DPA1*020105	3	10	5		
DPA1*020106	9	11	5		
DPA1*020201	12	11	5		
DPA1*020202	13	12	10	5	
DPA1*020203	14	5			
DPA1*0203	9	5			
DPA1*0301	15				
DPA1*0302	16				
DPA1*0401	17				

Table 15-2

Allele Number		Probe Number for Detection				
DPB1*010101	0	1				
DPB1*010102	2					
DPB1*020102	3	4	5	6	7	
DPB1*020103	8					
DPB1*020104	9					
DPB1*020105	10					
DPB1*020106	11					
DPB1*0202	12	13	5	14		
DPB1*030101	15	3	16	17		
DPB1*030102	18					
DPB1*0401	19	6	7			
DPB1*0402	3	4	6	7		
DPB1*0501	12	20	13	6		
DPB1*0601	16	17	21	6		
DPB1*0801	3	4	5			
DPB1*0901	22	16	5			
DPB1*1001	22	3	4	5		
DPB1*110101	23					
DPB1*110102	24					
DPB1*1301	15	5	25			
DPB1*1401	22	3	16	17		
DPB1*1501	23	26				
DPB1*1601	3	4	5	6		
DPB1*1701	22	16	5	6		
DPB1*1801	3	4	27			
DPB1*1901	13	5	25			
DPB1*200101	16	17	6			
DPB1*200102	28					
DPB1*2101	15	12	13	5	6	
DPB1*2201	12	13	5	6		

Table 15-3

Allele Number	Probe Number for Detection				
DPB1#2301	3	6	7		
DPB1#2401	13	14			
DPB1#2501	15	3	4	17	
DPB1#260101	29				
DPB1#2701	15	6			
DPB1#2801	4	17	27		
DPB1#2901	16	17	21		
DPB1#3001	22	13	5	6	
DPB1#3101	30				
DPB1#3201	31				
DPB1#3301	5	6	7		
DPB1#3401	30	26			
DPB1#3501	22	3	16		
DPB1#3601	15	12	20	13	6
DPB1#3701	3	4	5		
DPB1#3801	32				
DPB1#3901	6	7			
DPB1#4001	27				
DPB1#4101	33	34			
DPB1#4401	12	17	21		
DPB1#4501	3	4	17		
DPB1#4601	16	5	14		
DPB1#4701	13	5	14		
DPB1#4801	12	4	7	14	
DPB1#4901	4	6	7		
DPB1#5001	3	16	17		
DPB1#5101	19	4	6	7	
DPB1#5201	15	3	17		
DPB1#5301	4	27			
DPB1#5401	22	13	5		

Table 15-4

Allele Number	Probe Number for Detection			
DPB1#5501	22	3	5	6
DPB1#5601	19	17		
DPB1#5701	3	16	17	
DPB1#5801	12	5	6	
DPB1#5901	4	17	6	7
DPB1#6001	35			
DPB1#6101N	36			
DPB1#6201	12	20	27	
DPB1#6301	12	6		
DPB1#6401N	16	17	21	6
DPB1#6601	22	19	6	7
DPB1#6701	22	3	17	
DPB1#6801	3	4		
DPB1#6901	16	37		
DPB1#7001	3	16	17	
DPB1#7101	3	5	6	7
DPB1#7201	17	6	7	
DPB1#7301	4	17	7	
DPB1#7401	23	26		
DPB1#7501	3	4	7	
DPB1#7601	22	16	17	
DPB1#7701	38			
DPB1#7801	39			
DPB1#7901	15	3	4	
DPB1#8001	16	14		
DPB1#8101	4	5	6	7
DPB1#8201	14	40		
DPB1#8301	33			
DPB1#8401	13	41		
DPB1#8501	15	42		

Table 15-5

Allele Number	Probe Number for Detection				
DPB1*8601	43	16	5	14	
DPB1*8701	15	3	17	6	
DPB1*8801	15	16	5		
DPB1*8901	6				
DPB1*9001	19				
DPB1*9101	16	17	6		
DPB1*9201	15	16	17		
DPB1*9301	15	3	4	5	6
DPB1*9601	44				



Table 16-1

Allele Number	Probe Number for Detection			
DPA1#010301	0	1	2	
DPA1#010302	3			
DPA1#0104	4			
DPA1#0106	5			
DPA1#0107	6			
DPA1#0108	4	7		
DPA1#020101	8	5	7	
DPA1#020102	5	7		
DPA1#020103	9	7		
DPA1#020104	10			
DPA1#020105	3	9	7	
DPA1#020106	8	11	7	
DPA1#020201	3	11	7	
DPA1#020202	8	3	9	7
DPA1#020203	12	7		
DPA1#0203	8	7		
DPA1#0301	13			
DPA1#0302	12			
DPA1#0401	14			

Table 16-2

Allele Number	Probe Number for Detection						
DPB1*010101	0	1	2				
DPB1*010102	3						
DPB1*020102	4	5	6	7	8	9	
DPB1*020103	10						
DPB1*020104	11						
DPB1*020105	12						
DPB1*020106	13						
DPB1*0202	14	15	7	16			
DPB1*030101	17	5	18	19			
DPB1*030102	20						
DPB1*0401	4	21	8	9			
DPB1*0402	4	5	6	8	9		
DPB1*0501	4	14	22	15	8		
DPB1*0601	18	19	7	8			
DPB1*0801	5	6	7				
DPB1*0901	23	18	7				
DPB1*1001	23	6	7				
DPB1*110101	17	24					
DPB1*110102	25						
DPB1*1301	17	7	26				
DPB1*1401	23	5	18	19			
DPB1*1501	24	16					
DPB1*1601	4	5	6	7	8		
DPB1*1701	23	18	7	8			
DPB1*1801	5	6	27				
DPB1*1901	4	15	7	26			
DPB1*200101	18	19	8				
DPB1*200102	18	19	8				

Table 16-3

Allele Number	Probe Number for Detection				
DPB1*2101	17	14	15	7	8
DPB1*2201	4	14	15	7	8
DPB1*2301	4	5	8	9	
DPB1*2401	15	16			
DPB1*2501	17	5	6	19	
DPB1*260101	28				
DPB1*260102	17				
DPB1*2701	17	8			
DPB1*2801	6	19	27		
DPB1*2901	18	19	7		
DPB1*3001	23	29	15	7	8
DPB1*3101	30				
DPB1*3201	31				
DPB1*3301	4	7	8	9	
DPB1*3401	30	16			
DPB1*3501	23	5	18		
DPB1*3601	17	14	22	15	8
DPB1*3701	17	5	6	7	
DPB1*3801	32				
DPB1*3901	4	8	9		
DPB1*4001	4	27			
DPB1*4101	33	7			
DPB1*4401	14	19	7		
DPB1*4501	29	5	6	19	
DPB1*4601	4	18	7	16	
DPB1*4701	15	7	16		
DPB1*4801	14	6	9	16	
DPB1*4901	6	8	9		
DPB1*5001	5	18	19		
DPB1*5101	4	21	6	8	9

Table 16-4

Allele Number	Probe Number for Detection				
DPB1*5201	17	5	19		
DPB1*5301	4	6	27		
DPB1*5401	23	29	15	7	
DPB1*5501	23	7	8		
DPB1*5601	17	21	19		
DPB1*5701	5	18	19		
DPB1*5801	29	14	7	8	
DPB1*5901	6	19	8	9	
DPB1*6001	34				
DPB1*6101N	35				
DPB1*6201	14	22	27		
DPB1*6301	14	8			
DPB1*6401N	18	19	7	8	
DPB1*6501	4				
DPB1*6601	23	16			
DPB1*6701	23	5	19		
DPB1*6801	4	5	6		
DPB1*6901	18	36			
DPB1*7001	37	5	18	19	
DPB1*7101	5	7	8	9	
DPB1*7201	19	8	9		
DPB1*7301	6	19	9		
DPB1*7401	17	24	16		
DPB1*7501	5	6	9		
DPB1*7601	23	18	19		
DPB1*7701	38				
DPB1*7801	39				
DPB1*7901	17	5	6		
DPB1*8001	4	18	40		
DPB1*8101	4	6	7	8	9

Table 16-5

Allele Number	Probe Number for Detection				
	4	5	6	8	9
DPB1*8201	4	5	6	8	9
DPB1*8301	33				
DPB1*8401	41	42			
DPB1*8501	17	8			
DPB1*8601	23	7	16		
DPB1*8701	17	5	19	8	
DPB1*8801	17	18	7		
DPB1*8901	8				
DPB1*9001	21				
DPB1*9101	23	19	8		
DPB1*9201	17	18	19		
DPB1*9301	17	5	6	7	8
DPB1*9601	43				

(Example 9)

Probes for identification of HLA-DQ allele

Extraction of DNA from 1 ml of human blood was performed using GFX Genomic Blood DNA Purification  
5 Kit from Amersham Biosciences in the same manner as in Example 1.

Next, quantitative PCR was carried out in the same manner as in Example 1 except that probes in the probe lists DQ1A and DQ1B were used and 2  $\mu$ l of the  
10 mixed primers consisting of 1  $\mu$ l each of respective solutions of the following primers (10 pmol/ $\mu$ l) and 6  $\mu$ l of ultra pure water were used:

GGTGAGGTAAGTATCTTG (SEQ ID NO: 165)

TCCTTCTGGCTGTTCCAGTACTC (SEQ ID NO: 166).

15 After PCR amplification, referring to Amp Plot and Dissociation curves on a display of 5700 software, and to the allele-probe list (Table 19A, 19B-1 and 19B-2), it was identified as DQA1\*0103 and DQB1\*060101.

20 (Example 10)

Extraction of DNA from 1 ml of human blood was performed in the same way as in Example 3. PCR of human HLA-DQ was then performed in the same manner as in Example 2 except that 3  $\mu$ l of the mixed primer  
25 consisting of 1  $\mu$ l each of the solutions containing the following sequences at 10 pmol/ $\mu$ l respectively, and 12  $\mu$ l of ultra pure water were used:

GGTGAGGTAAGTATCTTG (SEQ ID NO: 165)

ATGATCCTAAACAAAGCTCTG (SEQ ID NO: 167)

TGTGCTACTTCACCAACGGGACG (SEQ ID NO: 168).

At the same time, a DNA microarray was prepared  
5 to identify the allele in the specimen described  
above in the same manner as in Example 2, except that  
probes in the probe list of Tables 18A, 18B-1 and  
18B-2 were used to form the probe spots respectively.

Then, hybridization was performed using the  
10 above specimen and the prepared DNA microarray in the  
same manner as in Example 2. Fluorometry measurement  
was conducted with GenePix4000B (Axon). Referring to  
the allele-probe list (Tables 20A, 20B-1 and 20B-2),  
it was identified as DQA1\*0103 and DQB1\*060101.

15

#### Allele list

DQA1\*010101

atgatcctaacaagcctgctgctgggggcccctgctctgaccaccgtgatgagccccctgtggagggtgaagaca  
tltgtggctgaccacgttgccctctgtgggtgtaacctgtaccagttttacggtccctctggccagtaacccatga  
20 atttgatggagatgagGagttctacgtggaccitggagaggaaggagactgcctggcgggtggcctgagttcagcaaa  
tttggagggtttgacccgcagggtgcactgagaaacatggctgtggcaaaacacaaacttgaacatcatgatataaac  
gtacaactctaccgtgctaccaatgaggttccctgagggtcacagtgtttccaagtcctccgtgacactgggtca  
gcccacaccctcatttgtctgtggacaacatctttccctctgtggcaacatcacatggctgagcaatgggcag  
tcagtcacagaagggtttctgagaccagcttccctctccaagagtgatcatctctcttcaagatcagttacctca  
25 ccttctctccctctgctgatgagattatgactgcaagggtggagcactggggcctggaccagcccttctgaaaca  
ctgggagccctgagattccagccccatgtctagagctcacagagactgtggctgcgccctgggggtgtctgtgggc  
ctcgtgggcattgtgggtgggcactgtcttcatcatccaaggcctgcgttcagttgggtgtctccagacaccaagggc

catttgtga (SEQ ID NO:169)

DQA1\*010102

atgatcctaacaagctctgctgctgggggcccccgctctgaccaccgtgatgagccccctgggaggatgaagaca  
ttgtggctgaccacgttgcctcttgggtgtaaacttgiaccagttttacgggtccccctggccagttacacccaatga  
5 atttgatggagatgaggagttctacgtggaccggagaggaaggagactgccctggcgggtggccctgagttcagcaaa  
tttggaggttttgaccgcagggtgcactgagaaacatggctgtggcaaaacacaacttgaacatcatgattaaac  
gctacaactctaccgtctgctaccaatgagggtccctgagggtcacagtggtttccaagctctcccgtagacactgggtca  
gccccacacccctcatttgccttgggacaacatcttccctccctgtgggtcaacatcacatggctgagcaatgggcag  
tcagtcacagaagggtgttctgagaccagcttccctccaagagtcatcttcttcaagatcagttacctca  
10 ccttccctcccttctgctgatgagatttatgactgcaagggtggagcacggggccctggaccagccctctctgaaaca  
ctgggagccctgagattccagccccatgtcagagctcacagagactgtggctctgcgccctggggtgtctgtgggc  
ctcgtgggcatttgggtgggcactgtcttcatcatccaaggccctgcgttcagttgggtgtctccagacaccaGgggc  
catttgtga (SEQ ID NO:170)

DQA1\*010201

15 atgatcctaacaagctctgctgctgggggcccccgctctgaccaccgtgatgagccccctgggaggatgaagaca  
ttgtggctgaccacgttgcctcttgggtgtaaacttgiaccagttttacgggtccccctggccagttacacccaatga  
atttgatggagatgagcagttctacgtggaccggagaggaaggagactgccctggcgggtggccctgagttcagcaaa  
tttggaggttttgaccgcagggtgcactgagaaacatggctgtggcaaaacacaacttgaacatcatgattaaac  
gctacaactctaccgtctgctaccaatgagggtccctgagggtcacagtggtttccaagctctcccgtagacactgggtca  
20 gccccacacccctcatttgccttgggacaacatcttccctccctgtgggtcaacatcacatggctgagcaatgggcag  
tcagtcacagaagggtgttctgagaccagcttccctccaagagtcatcttcttcaagatcagttacctca  
ccttccctcccttctgctgatgagatttatgactgcaagggtggagcacggggccctggaccagccctctctgaaaca  
ctgggagccctgagattccagccccatgtcagagctcacagagactgtggctctgtgccctggggtgtctgtgggc  
ctcAtgggcatttgggtgggcactgtcttcatcatccaaggccctgcgttcagttgggtgtctccagacaccaagggc  
25 catttgtga (SEQ ID NO:171)

DQA1\*010202

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5 gccaacacccctcatctgtcttgtggacaacatctttccctcttgggtcaacatcacatggctgagcaatgggcag  
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10 cattgtga (SEQ ID NO:172)

DQA1\*0103

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DQA1\*010401

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cattgtga (SEQ ID NO:174)

DQA1\*010402

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DQA1\*0105

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DQA1\*0106

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DQA1\*0201

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20 DQA1\*030101

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ccctgtga (SEQ ID NO:179)

## 5 DQA1\*0302

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## DQA1\*0303

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DQA1\*040101

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DQA1\*040102

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SEQ ID NO:183)

DQA1\*050101

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ccttgtga (SEQ ID NO:184)

DQA1\*050102

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DQA1\*0502

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Q ID NO:186)

DQA1\*0503

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DQA1\*0504

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DQA1\*0505

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20 DQA1\*060101

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5 DQA1\*060102

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ga (SEQ ID NO:191)

10 DQB1\*050101

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Q ID NO:192)

DQB1\*050102

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DQB1\*050201

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5 Q ID NO:194)

DQB1\*050202

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10 cgggatcctgcagagga (SEQ ID NO:195)

DQB1\*050301

gggccctgtgctacttcaccaacgggacggagcgcgtgcgggggttgaccagacacatctataaaccgagaggagtac  
gtgcgttctgacagcgacgtgggggtgtatcgggcggtagcggcgaggggcgccctAGcggcagtagctggaaca  
gccagaaggaagtcctggagggggcccgggcgtcggtaggacagAggtgtcagacacaactacgaggtggcgtaccg

15 cgggatcctgcagaggagatggagcccacagtaccatctcccatccaggacagaggccctcaaccaccacaac  
ctgctgatctgctcggtagacagatttctatccaagccagatcaaaagtcggtaggtttcggaatgatcaggaggaga  
cagccggcgttggtgccacccccctcatlaggaacggtagctggaccttccagatcctggtagctggaaatgac  
tccccagcgtggagatgtctacacctgccacgtggagcaccacagccctccagagcccatcaccgtggagtgg (SE  
Q ID NO:196)

20 DQB1\*050302

gacggagcgcgtgcgggggttgaccagacacatctataaaccgagaggagtacgtgcgttctgacagcgacgtgggg  
gtgtatcgggcggtagcggcgaggggcgccctAGtccgagtagctggaacagccagaaggaagtcctggag (SEQ  
ID NO:197)

DQB1\*0504

25 ggccctgtgctacttcaccaacgggacggagcgcgtgcgggggttgaccagatatactataaaccgagaagagtac  
gtgcgttctgacagcgacgtgggggtgtaccggcggtagcggcgaggggcgccctAGcggcagtagctggaaca  
gccagaaggacatcctggaggAggaccgggcgtcggtaggacagggtgtcagacacaact (SEQ ID NO:198)

DQB1\*0201

gggcatgtgctacttcaccaacgggacagagcgcgtgcgtcttgtgagcagaagcatctataaccgagaagagatc  
gtgcgcttcgacagcgacgtgggggagttccgggcggtagcgtgctggggctgccctgccgccgagttaciggaaca  
gccagaaggacatcctggagaggaaAcgggcggcggtaggacaggggtgtgcagacacaactaccagttggagctccg  
5 caccaccttgcagcggcgagtaggagcccacagtaccatctcccatccaggacagaggccctcaaccaccacaac  
ctgctggcttgcctggtagacagatttctatccagcccagatcaaagtcggtaggttctcggaatgaccaggaggaga  
cagctggcgttgtgtccaccccccttattaggaatggtagctggaccttccagatcctggtagtctggaaatgac  
tccccagcgtggagaCgtctacacctgccacgtggagcaccacagcctccagagccccatcacctggagttgg (SE  
Q ID NO:199)

10 DQB1\*0202

gggcatgtgctacttcaccaacgggacagagcgcgtgcgtcttgtgagcagaagcatctataaccgagaagagatc  
gtgcgcttcgacagcgacgtgggggagttccgggcggtagcgtgctggggctgccctgccgccgagttaciggaaca  
gccagaaggacatcctggagaggaaacgggcggcggtaggacaggggtgtgcagacacaactaccagttggagctccg  
caccaccttgcagcggcgagtaggagcccacagtaccatctcccatccaggacagaggccctcaaccaccacaac  
15 ctgctggcttgcctggtagacagatttctatccagcccagatcaaagtcggtaggttctcggaatgGccaggaggaga  
cagctggcgttgtgtccaccccccttattaggaatggtagctggaccttccagatcctggtagtctggaaatgac  
tccccagcgtggagaCgtctacacctgccacgtggagcaccacagcctccagagccccatcacctggagttgg (SE  
Q ID NO:200)

DQB1\*0203

gggcatgtgctacttcaccaacgggacagagcgcgtgcgtcttgtgagcagaagcatctataaccgagaagagatc  
gtgcgcttcgacagcgacgtgggggagttccgggcggtagcgtgctggggctgccctgAcgccgagttaciggaaca  
gccagaaggacatcctggagaggaaacgggcggcggtaggacaggggtgtgcagacacaactaccagttggagctccg  
caccaccttgcagcggcgaccccatccaggacagaggccctcaaccaccacaaccgtgctggcttgcctggtagag  
atttctatccagcccagatcaaagtcggtaggttctcggaatgGccaggaggagacagctggcgttgtgtccacccc  
25 ccttattaggaatggtagctggaccttccagatcctggtagtctggaaatgactccccagcgtggaga (SEQ ID  
NO:201)

DQB1\*030101

ggccatgtgctacttcaccaacgggacggagcgcgtgcgttatgtgaccagatacatctataaccgagaggagtlac  
gcacgcttcgacagcgacgtggAggtgtaccggcggtgacgccgctggggccgccIgAcgccgagtlactggaaca  
gccagaaggaagtcctggagaggacccgggcggagtggacacgggtgagacacacaactaccagttggagctccg  
cacgaccttgcagcggcgagtggagcccacagtgaccatctcccatccaggacagaggccctcaaccaccacaac  
5 ctgctggctcgtcagtgacagatttctatccagcccagatcaaagtcgggtggtttcggaatgaccaggaggaga  
caaccggcgttgtgtccaccccccttattaggaacgggtgactggaccttccagatcctgggtgagctggaaatgac  
tccccagcatggagaCgltctacacctgccacgtggagcaccacagcctccagaAccccatcacctggagtgg (SE  
Q ID NO:202)

DQB1\*030102

10 ggccatgtgctacttcaccaacgggacggagcgcgtgcgttatgtgaccagatacatctataaccgagaggagtlac  
gcgcgcttcgacagcgacgtggAggtgtaccggcggtgacgccgctggggccgccIgAcgccgagtlactggaaca  
gccagaaggaagtcctggagaggacccgggcggagtggacacgggtgagacacacaactaccagttggagctccg  
cacgaccttgcagcggcgag (SEQ ID NO:203)

DQB1\*0302

15 gggcatgtgctacttcaccaacgggacggagcgcgtgcgtctgtgaccagatacatctataaccgagaggagtlac  
gcAcgcttcgacagcgacgtgggggtgtatcggcggtgacgccgctggggccgccIgCgccgagtlactggaaca  
gccagaaggaagtcctggagaggacccgggcggagTggacacgggtgagacacacaactaccagttggagctccg  
cacgaccttgcagcggcgagtggagcccacagtgaccatctcccatccaggacagaggccctcaaccaccacaac  
ctgctggctcgtcagtgacagatttctatccagcccagatcaaagtcgggtggtttcggaatgaccaggaggaga  
20 caactggcgttgtgtccaccccccttattaggaacgggtgactggaccttccagatcctgggtgagctggaaatgac  
tccccagcgtggagacgltctacacctgccacgtggagcaccacagcctccagaaccccatcaTcgtggagtgg (SE  
Q ID NO:204)

DQB1\*030302

25 gggcatgtgctacttcaccaacgggacggagcgcgtgcgtctgtgaccagatacatctataaccgagaggagtlac  
gcacgcttcgacagcgacgtgggggtgtatcggcggtgacgccgctggggccgccIgAcgccgagtlactggaaca  
gccagaaggaagtcctggagaggacccgggcggagTggacacgggtgagacacacaactaccagttggagctccg  
cacgaccttgcagcggcgagtggagcccacagtgaccatctcccatccaggacagaggccctcaaccaccacaac

ctgctggctcgtcagtgacagatttctatccagcccagatcaaagtcgggtgggttcggaatgaccaggaggaga  
caactggcgltgtgtccaccccccttattaggaacggtagctggaccttccagatcctggtagtgcctggaaatgac  
tccccagcgtggagacgtctacacctgccacgtggagcaccacagcctccagaaccccatcaTcgtggagtgg (SE  
Q ID NO:205)

5 DQB1\*030303

gggcatgtgtctacttcaccaacgggacggagcgcgtgcgtTtgtgaccagatacatctataaaccgagaggagtac  
gcgcgcttcgacagcgacgtgggggtgtatcgggcggtagacccgctggggcCgcctgAcgccgagtagctggaaca  
gccagaaggaagtcctggagAggacccgggaggagTtggacacggtagtcagacacaactaccagtggagctccg  
cacgaccttcagcggcgag (SEQ ID NO:206)

10 DQB1\*0304

ggccatgtgtctacttcaccaacgggacggagcgcgtgcgtTtgtgaccagatacatctataaaccgagaggagtac  
gcacgcttcgacagcgacgtggAggtgtaccgggaggtagacccgctggggccgctgCgcccagtagctggaaca  
gccagaaggaagtcctggagaggacccgggaggagtggacacggtagtcagacacaactaccagtggagctccg  
cacgaccttcagcggcgagtaggagcccacagtaccatctccccatccaggacagaggccctcaaccaccacaac

15 ctgctggctcgtcagtgacagatttctatccagcccagatcaaagtcgggtgggttcggaatgaccaggaggaga  
caaccggcgltgtgtccaccccccttattaggaacggtagctggaccttccagatcctggtagtgcctggaaatgac  
tccccagcatggagaCgtctacacctgccacgtggagcaccacagcctccagaAccccatcacctggagtgg (SE  
Q ID NO:207)

DQB1\*030501

20 gggcatgtgtctacttcaccaacgggacCgagcgcgtgcgggtgtgaccagatacatctataaaccgagaggagtac  
gcgcgcttcgacagcgacgtgggggtgtatcgggcggtagacccgctggggccgctgccgcccagtagctggaaca  
gccagaaggaagtcctggagaggacccgggaggagTtggacacggtagtcagacacaactaccagtggagctccg  
cacgaccttcagcggcgagtaggagcccacagtaccatctccccatccaggacagaggccctcaaccaccacaac  
ctgctggctcgtcagtgacagatttctatccagcccagatcaaagtcgggtgggttcggaatgaccaggaggaga  
25 caactggcgltgtgtccaccccccttattaggaacggtagctggaccttccagatcctggtagtgcctggaaatgac  
tccccagcgtggagacgtctacacctgccacgtggagcaccacagcctccagaaccccatcatcgtggagtgg (SE  
Q ID NO:208)

DQB1\*030502

gggcatgtgctacttcaccaacgggacggagcgcgtgcgggtgtgaccagatacatctataaccgagaggagtag  
gcgcgcttcgacagcgacgtgggggtgtatcgggcgggtgacccgcctggggccgctgCgcccagtagtggaaaca  
gccagaaggaagtcctggagAggacccgggaggagtaggacaCgggtgtgcagacacaactaccagtaggagctccg  
5 cagaccttgcagcggcgag (SEQ ID NO:209)

DQB1\*0306

gggcatgtgctacttcaccaacgggacggagcgcgtgcgtcttgtgaccagatacatctataaccgagaggagtag  
gcacgcttcgacagcgacgtgggggtgtatcgggcgggtgacccgcctggggcCgctgacgccgagtagtggaaata  
gccagaaggacatcctggaggaggaccggggtcggtaggacaccgtAtgcagacacaactaccagtaggagctccg  
10 cagaccttgcagcggcgag (SEQ ID NO:210)

DQB1\*0307

gggcatgtgctacttcaccaacgggacggagcgcgtgcgtcttgtgaccagatacatctataaccgagaggagtag  
gcacgcttcgacagcgacgtgggggtgtatcgggTggtagccgcctggggccgctgcccggagtagtggaaaca  
gccagaaggaagtcctggagaggaccgggaggagtaggacacgggtgtgcagacacaactaccagtaggagctccg  
15 cagaccttgcagcggcga (SEQ ID NO:211)

DQB1\*0308

gggcatgtgctacttcaccaacgggacggagcgcgtgcgtcttgtgaccagatacatctataaccgagaggagtag  
gcAcgcttcgacagcgacgtgggggtgtatcgggcgggtgacccgcctggggccgctgCgcccagtagtggaaaca  
gccagaaggaagtcctggaggggaccgggaggagtaggacaCgggtgtgcagacacaactaccagtaggagctccg  
20 cagaccttgcagcggcgag (SEQ ID NO:212)

DQB1\*0309

ggccatgtgctacttcaccaacgggacggagcgcgtgcgttattgtgaccagatacatctataaccgagaggagtag  
gcacgcttcgacagcgacgtggagggtgacgggaggtagccgcctggggccgctgacgccgagtagtggaaaca  
gccagaaggaagtcctggagaggaccgggaggagtaggacacgggtgtgcagacacaactaccagtaggagctccg  
25 cagaccttgcagcggcgagtaggagcccacagtaccatctcccatccaggacagaggccctcaaccaccacaac  
ctgctgggtcgtcagtagacagattctatccagcccagatcaaagtccggtgggttcggaatgaccaggaggaga  
caaccggcgttgtgtccaccccccttattaggaacggtagctggacctccagatccctgggtgatgctggaaatgac

tccccagcaigcC...gcttacacctgccacgtggagcacccagcctccagaaccccatcacctggagtgg (SEQ ID NO:213)

DQB1\*0310

ggccatgtgctacttcaccaacgggacggagcgcgtgcgttatgtgaccagatacatctataaccgagaggagttac  
5 gcacgcttcgacagcgacgtgggggtgtatcgggcgggtgacgccgtggggccgctgAcgccgagttactggaaca  
gccagaaggaagtcctggagaggacccgggcggagtggacacgggtgtcagacacaactaccagttggagctccg  
cacgaccttcagcggcgagttggagcccacagtgaccatctcccatccaggacagaggccctcaaccaccacaac  
ctgctgggtcgtcagtgacagattctatccagcccagatcaaatccggtgggtttcggaatgaccaggaggaga  
caaccggcgttgggtccaccccccttattaggaacgggtgactggaccttcagatcctgggtgactggaaatgac  
10 tccccagcAtggagaCgtctacacctgccacgtggagcacccagcctccagaAccccatcacctggagtgg (SEQ  
ID NO:214)

DQB1\*0311

gggcctgtgctacttcaccaacgggacggagcgcgtgcgtctgtgaccagatacatctataaccgagaggagttac  
gcAcgcttcgacagcgacgtgggggtgtatcgggcgggtgacgccgtggggccgctgCgccgagttactggaaca  
15 gccagaaggaagtcctggagAggacccgggcggagtggacaCgggtgtcagacacaactaccagttggagctccg  
cacgaccttcagcggcgag (SEQ ID NO:215)

DQB1\*0312

ggccatgtgctacttcaccaacgggacggagcgcgtgcgtTgtgaccagatacatctataaccgagaggagttac  
gcAcgcttcgacagcgacgtgggggtgtatcgggcgggtgacgccgtggggccgctgAcgccgagttactggaaca  
20 gccagaaggaagtcctggagAggacccgggcggagTggacacgggtgtcagacacaactaccagttggagctccg  
cacgaccttcagcggcgag (SEQ ID NO:216)

DQB1\*0313

ggccatgtgctacttcaccaacgggacggagcgcgtgcgttatgtgaccagatacatctataaccgagaggagttac  
gcacgcttcgacagcgacgtggagggtgtaccgggcgggtgacgccgtggggccgctgacgccgagttactggaaca  
25 gccagaaggaagAccttggagaggacccgggcggagtggacacgggtgtcagacacaactaccagttggagctccg  
cacgaccttcagcggcgag (SEQ ID NO:217)

DQB1\*0401

gggcatgtgctacttcaccaacgggaccgagcTcgtgcgggtgtgaccagatacatctataaccgagaggagtac  
gcgcgcttcgacagcgacgtgggggtgtatcgggcggtagcccgctggggcggcttgacgccgagtagtgaata  
gccagaaggacatccctggaggaggaccgggcgtcggtagacaccgtatgcagacacaactaccagttggagctccg  
cacgaccttgcagcggcgagtaggagcccacagtaccatctcccatccaggacagaggccctcaaccaccacaac  
5 ctgctggcttgcctcagtagacagatttctatccagcccagatcaaagtcgggtggtttcggaatgaccaggaggaga  
caactggcgttgtgtccaccccccttattaggaacggtagctggaccttccagatccctggtagtctggaaatgac  
tccccagcgtggagacgtctacacctgccacgtggagcaccacagccctccagaaccccatcatcgtggagtgg (SE  
Q ID NO:218)

DQB1\*0402

10 gggcatgtgctacttcaccaacgggaccgagcgctgcgggtgtgaccagatacatctataaccgagaggagtac  
gcgcgcttcgacagcgacgtgggggtgtatcgggcggtagcccgctggggcggcTtagcccgagtagtgaata  
gccagaaggacatccctggaggaggaccgggcgtcggtagacaccgtatgcagacacaactaccagttggagctccg  
cacgaccttgcagcggcgagtaggagcccacagtaccatctcccatccaggacagaggccctcaaccaccacaac  
ctgctggcttgcctcagtagacagatttctatccagcccagatcaaagtcgggtggtttcggaatgaccaggaggaga  
15 caactggcgttgtgtccaccccccttattaggaacggtagctggaccttccagatccctggtagtctggaaatgac  
tccccagcgtggagacgtctacacctgccacgtggagcaccacagccctccagaaccccatcatcgtggagtgg (SE  
Q ID NO:219)

DQB1\*060101

ggccatgtgctacttcaccaaTgggacggagcgctgcgttatgtgaccagatacatctataaccgagaggaggac  
20 gtgcgcttcgacagcgacgtgggggtgtatcgggcggtagcccgaggggcggccttagcccgagtagtgaaca  
gccagaaggacatccctggagaggaccgagcggagtaggacacgggtgtgcagacacaactaccaggtggcgttccg  
cgggatcttgcagaggagagtggagcccacagtaccatctcccatccaggacagaggccctcaaccaccacaac  
ctgctggcttgcctcggtagacagatttctatccagggccagatcaaagtcgggtggtttcggaatgaccaggaggaga  
cagctggcgttgtgtccaccccccttattaggaacggtagctggaccttccagatccctggtagtctggaaatgac  
25 tccccagcatggagacgtctacacctgccacgtggagcaccacagccctccagagcccatcaccgtggagtgg (SE  
Q ID NO:220)

DQB1\*060102

gccatgtgctacttcaccaacgggacggagcgcgtgcgttaigtgaccagatacatctataaaccgagaggaggacg  
tgcgccttcgacagcgacgtgggggtgtatcgggcgggtgacCccgcagggcgccctgacgccgagtagtggaaacg  
ccagaaggacatccctggagaggacccgagcggagttggacacgggtgtgcaga (SEQ ID NO:221)

DQB1\*060103

5 ggccatgtgctacttcaccaatgggacggagcgcgtgcgttaigtgaccagatacatctataaaccgagaggaggac  
gtgcgccttcgacagcgacgtgggggtgtatcgggcgggtgacgccgcagggcgccctgacgccgagtagtggaaacg  
gccagaaggacatccctggagaggacccgagcggagttggacacgggtgtgcagacacaaactacgaggtggcgttccg  
cgggatcttgcagaggagagtgagagccacagtaccatctccccatccaggacagaggccctcaaccaccacaac  
ctgctgggtctgctcgggtgacagatttctatccaggccagatcaaagtcgggtgggttccggaatgaccaggaAgaga  
10 cagctggcgttgtgtccaccccccttattaggaacgggtgactggacctccagatccctgggtgatgctggaaatgac  
tccccagcatggagacgtctacacctgccacgtggagcaccacagctccagagccccatcacctggagttgg (SE  
Q ID NO:222)

DQB1\*0602

gggcatgtgctacttcaccaacgggacggagcgcgtgcgtcttgtgaccagaTaatctataaaccgagaggagtag  
15 gcgcgccttcgacagcgacgtgggggtgtaccgcgcgggtgacgccgcagggcgccctgatgccgagtagtggaaacg  
gccagaaggaagtcctggaggggacccgggcggagttggacacgggtgtgcagacacaaactacgaggtggcgttccg  
cgggatcttgcagaggagagtgagagccacagtaccatctccccatccaggacagaggccctcaaccaccacaac  
ctgctgggtctgctcgggtgacagatttctatccaggccagatcaaagtcgggtgggttccggaatgatcaggaggaga  
cagccggcgttgtgtccaccccccttattaggaatgggtgactggacTtccagatccctgggtgatgctggaaatgac  
20 tccccagcgtggagatgtctacacctgccacgtggagcaccacagctccagagccccatcacctggagttgg (SE  
Q ID NO:223)

DQB1\*0603

gggcatgtgctacttcaccaacgggacggagcgcgtgcgtcttgtAaccagacacatctataaaccgagaggagtag  
gcgcgccttcgacagcgacgtgggggtgtaccgcgcgggtgacgccgcagggcgccctgatgccgagtagtggaaacg  
25 gccagaaggaagtcctggaggggacccgggcggagttggacacgggtgtgcagacacaaactacgaggtggcgttccg  
cgggatcttgcagaggagagtgagagccacagtaccatctccccatccaggacagaggccctcaaccaccacaac  
ctgctgggtctgctcgggtgacagatttctatccaggccagatcaaagtcgggtgggttccggaatgatcaggaggaga



cagccggcggtgtgtccaccccccttattaggaatggtagctggacTttccagatcctggtagcttgaaatgac  
tccccagcgtggagatgtctacacctgccacgtggagcaccacagcctccagagccccatcacctggagtg (SE  
Q ID NO:224)

DQB1\*060401

- 5 gggcatgtgctacttcaccaacgggacggagcgcgtgcgtcttgtAaccagacacatctataaccgagaggagtac  
gcgcgttcgacagcgacgtgggggtgtaccggcggtgacgccgcagggcgccctgttgccgagtactggaaca  
gccagaaggaagtcctggagAggacccggcgagttggacacggtgtgcagacacaactacgaggtgggtaccg  
cgggatcctgcagaggagagtgaggccacagtaccatctccccatccaggacagagggcccaaccaccacaac  
ctgtcgtgtgtcgtcgtgacagatttctatccaggccagatcaaagtcAggtgttccgaatgatcaggaggaga  
10 cagccggcggtgtgtccaccccccttattaggaatggtagctggacTttccagatcctggtagcttgaaatgac  
tccccagcgtggagatgtctacacctgccacgtggagcaccacagcctccagagccccatcacctggagtg (SE  
Q ID NO:225)

DQB1\*060402

- gggcatgtgctacttcaccaacgggacggagcgcgtgcgtcttgtAaccagacacatctataaccgagaggagtac  
15 gcgcgttcgacagcgacgtgggggtgtaccgCgcgtgacgccgcagggcgccctgttgccgagtactggaaca  
gccagaaggaagtcctggagAggAcggcgcgagttggacacggtgtgcagacacaactacgaggtggGgtaccg  
cgggatcctgcagaggagagtgaggccacagtaccatctccccatccaggacagaggcc (SEQ ID NO:226)

DQB1\*060501

- gggcctgtgctacttcaccaacgggacggagcgcgtgcgtcttgtAaccagaTatatctataaccgagaggagtac  
20 gcgcgttcgacagcgacgtgggggtgtaccggcggtgacgccgcagggcgccctgttgccgagtactggaaca  
gccagaaggaagtcctggagAggAcggcgcgagttggacacggtgtgcagacacaactacgaggtggGgtaccg  
cgggatcctgcagaggagagtgaggccacagtaccatctccccatccaggacagaggcc (SEQ ID NO:227)

DQB1\*060502

- ggacggagcgcgtgcgtcttgtAaccagatatactataaccgagaggagtacgcgcgttcgacagcgacgtggg  
25 ggtgtaccggcggtgacgccgcagggcgccctgtCgccgagtactggaacagccagaaggaagtcctggagAgg  
AcggcgcgagttggacaCg (SEQ ID NO:228)

DQB1\*0606

ggacggagcgcgctgcgtcttgtAaccagaTacaTctataaaccgagaggagTaccgcgcttcgacagcgacgtggg  
ggtgtaccggcggtgacgccgcagggcgccctgttgccgagTactggaacagccagaaggaagtcctggagAgg  
AccggggcgcggtggacagggTg (SEQ ID NO:229)

DQB1\*0607

- 5 gggcatgtgctacttcaccaacgggacggagcgcgctgcgtcttgtAaccagacacatctataaaccgagaggagTacc  
gcgcgcttcgacagcgacgtgggggtgtaccgcggtgacgccgcagggcgccctgtAtgccgagTactggaaca  
gccagaaggaagtcctggagAggAccggggcgagTtggacacggtgtgcagacacaactacgaggtggGgtaccg  
cgggatcc (SEQ ID NO:230)

DQB1\*0608

- 10 gggcatgtgctacttcaccaacgggacggagcgcgctgcgtcttgtAaccagacacatctataaaccgagaggagTacc  
gcgcgcttcgacagcgacgtgggggtgtaccgcggtgacgccgcagggcgccctgttgccgagTactggaaca  
gccagaaggaagtcctggaggggacccgggcggagTtggacacggtgtgcagacacaactacgaggtggcgTccg  
cgggatcT (SEQ ID NO:231)

DQB1\*0609

- 15 gggcatgtgctacttcaccaacgggacggagcgcgctgcgtcttgtAaccagaTacaTctataaaccgagaggagTacc  
gcgcgcttcgacagcgacgtgggggtgtaccggcggtgacgccgcagggcgccctgttgccgagTactggaaca  
gccagaaggaagtcctggagAggacccgggcggagTtggacacggtgtgcagacacaactacgaggtgggTaccg  
cgggatccTgcagaggagagTggagcccacagTgacatctccccTccaggacagaggccccTcaaccaccacaac  
ctgctggTctgctcggtgacagatTtctatccaggccagatcaaagTccAgTggTttcggaatgatcaggaggaga

- 20 cagccggcgTtgtgtccacccccTatTtaggaatggTgactggactTtccagatccTggtgatgtcTggaaatgac  
TccccagcgTggagatgtctacacctgccacgtggagacccccagccTccagagccccatcacctggagTgg (SE  
Q ID NO:232)

DQB1\*0610

- 25 gggcatgtgctacttcaccaacgggacggagcgcgctgcgtcttgtAaccagatacatctataaaccgagaggagTacc  
gcgcgcttcgacagcgacgtgggggtgtaccgcggtgacgccgcagggcgccctAGcgccgagTactggaaca  
gccagaaggaagtcctggaggggacccgggcggagTtggacacggtgtgcagacacaactacgaggtggcgTccg  
cgggatcTtgcagaggagag (SEQ ID NO:233)

DQB1\*061101

gggcatgtgctacttcaccaacgggacggagcgcgtgcgtcttggaccagaTaatctataaaccgagaggagtiac  
gcgcgcttcgacagcgacgtgggggtgtaccgCgcggtgacgccgcaggggcggcctgAtgccgagtiactggaaca  
gccagaaggaagtcctggaggggacccgggaggagtggacacgggtgtgcagacacaactacgaggtaggcgttccg  
5 cgggatcTtgcagagg (SEQ ID NO:234)

DQB1\*061102

gggcatgtgctacttcaccaacgggacggagcgcgtgcgtcttgtAaccagaTaatctataaaccgagaggagtiac  
gcgcgcttcgacagcgacgtgggggtgtaccgCgcggtgacgccgcaggggcggcctgAtgccgagtiactggaaca  
gccagaaggaagtcctggaggggacccgggaggagtggacacgggtgtgcagacacaactacgaggtaggcgttccg  
10 cgggatcTtgcagaggagag (SEQ ID NO:235)

DQB1\*0612

gggcatgtgctacttcaccaacgggacggagcgcgtgcgtcttgtAaccagaTaatctataaaccgagaggagtiac  
gcgcgcttcgacagcgacgtgggggtgtaccggcggtgacgccgcaggggcggcctgtgccgagtiactggaaca  
gccagaaggaagtcctggaggggacccgggaggagtggacacgggtgtgcagacacaactacgaggtaggggtaccg  
15 cgggatcctgcagaggagagtggagcccacagtgaccatctcccatccaggacagaggccctcaaccaccacaac  
ctgcctggctgtctgggtgacagattctatccaggccagatcaaagtcAgtaggtttcggaatgatcaggaggaga  
cagccggcgttgtgtccacccccctattaggaatgggtgactggactttccagatccctgggtgatgtcggaaatgac  
tccccagcgtggagatgtctacacctgccacgtggagcaccacagctccagagcccatcaccgtggagtgg (SE  
Q ID NO:236)

20 DQB1\*0613

gggcatgtgctacttcaccaacgggacggagcgcgtgcgtcttggaccagaTaatctataaaccgagaggagtiac  
gcgcgcttcgacagcgacgtgggggtgtaccgCgcggtgacgccgcaggggcggcctgtgccgagtiactggaaca  
gccagaaggaagtcctggaggggacccgggaggagtggacacgggtgtgcagacacaactacgaggtaggcgtTccg  
cgggat (SEQ ID NO:237)

25 DQB1\*0614

gggcatgtgctacttcaccaacgggacggagcgcgtgcgtcttgtAaccagacacatctataaaccgagaggagtiac  
gcgcgcttcgacagcgacgtgggggtgtaccgCgcggtgacgccgcaggggcggcctgAtgccgagtiactggaaca

gccagaaggaagtcctggaggggacccgggcggagttggacacggtgtgcagacacaactacgaggtggcgttccg  
cgggatcTtgcagaggagag (SEQ ID NO:238)

DQB1\*0615

gggcaigtgtctacttcaccaacgggacggagcgcggtgcgtcttgtgaccagaTaatctataaccgagaggagttac  
5 ggcgcgttcgacagcgacgtgggggtgtaccgcgcggtgacgccgcaggggcggcctgAtgccgagttactggaaca  
gccagaaggaagtcctggagAggAcccgggcggagttggacacggtgtgcagacacaactacgaggtggGgtaccg  
cgggatcctgcagaggagag (SEQ ID NO:239)

DQB1\*0616

gggcaigtgtctacttcaccaacgggacggagcgcggtgcgtcttgtgaccagaTaatctataaccgagaggagttac  
10 ggcgcgttcgacagcgacgtgggggtgtaccgcgcggtgacgccgcaggggcggcctgAtgccgagAactggaaca  
gccagaaggaagtcctggaggggacccgggcggagttggacacggtgtgcagacacaactacgaggtggcgttccg  
cgggatcttgcagaggagag (SEQ ID NO:240)

DQB1\*0617

gggcaigtgtctacttcaccaacgggacggagcgcggtgcgtcttgtAaccagacacatctataaccgagaggagttac  
15 ggcgcgttcgacagcgacgtgggggtgtaccgggcgggtgacgccgcaggggcggcctgttgccgagttactggaaca  
gccagaaggaagtcctggagggggcccgggcggagttggacacggtgtgcagacacaactacgaggtggGgtaccg  
c (SEQ ID NO:241)

DQB1\*0618

gggcaigtgtctacttcaccaacgggacggagcgcggtgcgtcttgtAaccagatacatctataaccgagaggagttac  
20 ggcgcgttcgacagcgacgtgggggtgtaccgggcgggtgacgccgcaggggcggcctgttgccgagttactggaaca  
gccagaaggaagtcctggagAggacccgggcggagttggacacggtgtgcagacacaactacgaggtggcgttccg  
cgggatcTtgcagaggag (SEQ ID NO:242)

DQB1\*0619

gggcaigtgtctacttcaccaacgggacggagcgcggtgcgtctTtgtgaccagaTaatctataaccgagaggagttac  
25 ggcgcgttcgacagcgacgtgggggtgtatcgggcggtgacgccgcTggggcggcctgAtgccgagttactggaaca  
gccagaaggaagtcctggaggggacccgggcggagTtggacacggtgtgcagacacaactacgaggtggcgttccg  
cgggatcTtgcagaggagag (SEQ ID NO:243)

DQB1\*0620

gggcctgtgctacitcaccaacgggacggagcgcgtgcgtcttgigaccagaTacaatcctaaaccgagaggagtag  
gcgcgcttcgacagcgacgtgggggtgtaccgCgcggtagcgcgcagggcgccctgAtgccgagtagtgaaca  
gccagaaggaagtcctggaggggacccgggcggagtggacacgggtgtgcagacacaactacgaggtaggcgtTccg

5 c (SEQ ID NO:244)

In the following, Probe Lists DQ1 and DQ2 are  
shown in Tables 17A, 17B-1 and 17B-2 and tables 18A,  
18B-1 and 18B-2 respectively. Tables 19A, 19B-1 and  
10 19B-2 and Tables 20A, 20B-1 and 20B-2 show Allele-  
Prove Lists.

Table 17A

Probe No.	Base Sequence
0	t gaa ttt gat gga gat gag G (SEQ ID No: 1)
1	ggt gct tcc aga cac caG (SEQ ID No: 2)
2	gg ttg tct gtg ggc ctc A (SEQ ID No: 3)
3	cag ccc aac acc ctc atC (SEQ ID No: 4)
4	g ctg agc aat ggg cac G (SEQ ID No: 5)
5	ca gag act gtg gtc tgc A (SEQ ID No: 6)
6	c cct tgt gga ggt gaa gG (SEQ ID No: 7)
7	cct gtg gtc aac atc acC (SEQ ID No: 8)
8	ccc tgt gga ggt gaa gG (SEQ ID No: 9)
9	c ctg gag agg aag gag G (SEQ ID No: 10)
10	tg cct ctg ttc cac aga C (SEQ ID No: 11)
11	x ag cct gag att cca A (SEQ ID No: 12)
12	gcc ctg acc acc gtg aC (SEQ ID No: 13)
13	c acc ttc ctc cct tct gA (SEQ ID No: 14)
14	tt aaa cgc tcc aac tct acT (SEQ ID No: 15)
15	cc aga cac caa ggg ccC (SEQ ID No: 16)
16	ca gtg ttt tcc aag tct ccT (SEQ ID No: 17)
17	g cac tgg ggc ctg gac A (SEQ ID No: 18)
18	g gtc tgc gcc ctg ggA (SEQ ID No: 19)
19	ct gac cac gtt gcc tct tA (SEQ ID No: 20)
20	c cta aaa cat aac ttg aac agT (SEQ ID No: 21)
21	c aga caa ttt aga ttt gac cG (SEQ ID No: 22)
22	tc acc ctc ctc cct tct T (SEQ ID No: 23)
23	tg tac cag tct tac ggt cT (SEQ ID No: 24)
24	ag gtg gag cac tgg ggA (SEQ ID No: 25)
25	ggt ccc tct ggc cag tT (SEQ ID No: 26)
26	cc aag tct ccc gtg acG (SEQ ID No: 27)
27	gca ctg aca aac atc gcC (SEQ ID No: 28)

Table 17B-1

Probe No.	Base Sequence
0	g ggg gtg tac cgg gcA ( SEQ ID No: 29)
1	cg cag ggg cgg cct gT ( SEQ ID No: 30)
2	ag ggg gcc cgg gcg T ( SEQ ID No: 31)
3	gg gcg tcg gtg gac aG ( SEQ ID No: 32)
4	gg gcg tcg gtg gac agA ( SEQ ID No: 33)
5	ca gat ttc tat cca agc caC ( SEQ ID No: 34)
6	gc gac gtg ggg gtg taT ( SEQ ID No: 35)
7	cg cag ggg cgg cct aG ( SEQ ID No: 36)
8	g cag ggg cgg cct agC ( SEQ ID No: 37)
9	cg cag ggg cgg cct gA ( SEQ ID No: 38)
10	g cag ggg cgg cct gaC ( SEQ ID No: 39)
11	g aag gac atc ctg gag gA ( SEQ ID No: 40)
12	g gac atc ctg gag agg aaA ( SEQ ID No: 41)
13	ct ccc cag cgt gga gaC ( SEQ ID No: 42)
14	c cgg tgg ttt cgg aat gG ( SEQ ID No: 43)
15	ctg ctg ggg ctg cct gA ( SEQ ID No: 44)
16	c ttc gac agc gac gtg gA ( SEQ ID No: 45)
17	cg ctg ggg ccg cct gA ( SEQ ID No: 46)
18	ct ccc cag cat gga gaC ( SEQ ID No: 47)
19	cac ccc agc ctc cag aA ( SEQ ID No: 48)
20	aac cga gag gag tac gcA ( SEQ ID No: 49)
21	g ctg ggg ccg cct gC ( SEQ ID No: 50)
22	agg acc cgg gcg gag T ( SEQ ID No: 51)
23	c ctc cag aac ccc atc aT ( SEQ ID No: 52)
24	cg gag cgc gtg cgt cT ( SEQ ID No: 53)
25	g acg ccg ctg ggg cC ( SEQ ID No: 54)
26	cag aag gaa gtc ctg gag A ( SEQ ID No: 55)
27	tac ttc acc aac ggg acC ( SEQ ID No: 56)

Table 17B-2

Probe No.	Base Sequence
28	cgg gcg gag ttg gac aC (SEQ ID No: 57)
29	cg tcg gtg gac acc gtA (SEQ ID No: 58)
30	gtg ggg gtg tat cgg gT (SEQ ID No: 59)
31	tg act ccc cag cat gcC (SEQ ID No: 60)
32	g gaa atg act ccc cag cA (SEQ ID No: 61)
33	gg aac agc cag aag gaa gA (SEQ ID No: 62)
34	acc aac ggg acc gag cT (SEQ ID No: 63)
35	g ccg ctg ggg cgg cT (SEQ ID No: 64)
36	cc atg tgc tac ttc acc aaT (SEQ ID No: 65)
37	tg tat cgg gcg gtg acC (SEQ ID No: 66)
38	g ttt cgg aat gac cag gaA (SEQ ID No: 67)
39	gtg cgt ctt gtg acc aga T (SEQ ID No: 68)
40	g gcg ttc cgc ggg atc T (SEQ ID No: 69)
41	t agg aat ggt gac tgg acT (SEQ ID No: 70)
42	gag cgc gtg cgt ctt gtA (SEQ ID No: 71)
43	ca ggc cag atc aaa gtc cA (SEQ ID No: 72)
44	c gtg ggg gtg tac cgC (SEQ ID No: 73)
45	ag gaa gtc ctg gag agg A (SEQ ID No: 74)
46	a cac aac tac gag gtg gG (SEQ ID No: 75)
47	gtg cgt ctt gta acc aga T (SEQ ID No: 76)
48	g cag ggg cgg cct gtC (SEQ ID No: 77)
49	c aac tac gag gtg gcg tT (SEQ ID No: 78)
50	g cgg cct gat gcc gag A (SEQ ID No: 79)
51	gg gcg gtg acg ccg cT (SEQ ID No: 80)
52	cg ctg ggg cgg cct gA (SEQ ID No: 81)
53	ggg acc cgg gcg gag T (SEQ ID No: 82)



Table 18A

Probe No.	Base Sequence
0	gga gat gag gag ttc tac g (SEQ ID No: 83)
1	c aga cac caG ggg' cca tt (SEQ ID No: 84)
2	gtg ggc ctc Atg ggc att (SEQ ID No: 85)
3	c acc ctc atC tgt ctt gtg (SEQ ID No: 86)
4	aat ggg cac Gca gtc aca (SEQ ID No: 87)
5	g gtc tgc Acc ctg ggg (SEQ ID No: 88)
6	ga ggt gaa gGc att gtg g (SEQ ID No: 89)
7	c aac atc acC tgg ctg ag (SEQ ID No: 90)
8	gg aag gag Gct ggc tgg (SEQ ID No: 91)
9	ctg ttc cac aga Ctt aga c c ttt (SEQ ID No: 92)
10	gag att cca Aca cct atg tc (SEQ ID No: 93)
11	c acc gtg aCg agc cct t (SEQ ID No: 94)
12	ctc cct tct gAt gat gag at (SEQ ID No: 95)
13	c aac tct acI gct gct acc (SEQ ID No: 96)
14	c atc atc cGa ggc ctg c (SEQ ID No: 97)
15	c aag tct ccT gtg acg ct (SEQ ID No: 98)
16	ggc ctg gac Aag cct ctt (SEQ ID No: 99)
17	c gcc ctg ggA ttg tct gt (SEQ ID No: 100)
18	gtt gcc tct tAt ggt gta aa (SEQ ID No: 101)
19	aac ttg aac agT ctg att aaa c (SEQ ID No: 102)
20	a cg ttt gac cGg caa ttt gca c (SEQ ID No: 103)
21	ctc cct tct Tct gag gag (SEQ ID No: 104)
22	ct tac ggt cTc tct ggc c (SEQ ID No: 105)
23	g cac tgg ggA ctg gac aa (SEQ ID No: 106)
24	ct ggc cag tTc acc cat g (SEQ ID No: 107)
25	ccc gtg acG ctg ggt c (SEQ ID No: 108)
26	ca aac atc gcC gtg aca aaa (SEQ ID No: 109)

Table 18B-1

Probe No.	Base Sequence
0	tac cgg gcA gtg acg cc (SEQ ID No: 110)
1	g cgg cct gTt gcc gag (SEQ ID No: 111)
2	c cgg gcg Tcg gtg gac (SEQ ID No: 112)
3	g gtg gac aGg gtg tgc a (SEQ ID No: 113)
4	g gtg gac agA gtg tgc ag (SEQ ID No: 114)
5	t cca agc caC atc aaa gtc (SEQ ID No: 115)
6	ggg gtg taT cgg gcg g (SEQ ID No: 116)
7	g cgg cct aGc gcc gag (SEQ ID No: 117)
8	cgg cct agC gcc gag t (SEQ ID No: 118)
9	g cgg cct gAc gcc gag (SEQ ID No: 119)
10	cgg cct gaC gcc gag t (SEQ ID No: 120)
11	g cgg cct gAt gcc gag (SEQ ID No: 121)
12	c ctg gag gAg gac cgg (SEQ ID No: 122)
13	gag agg aaA cgg gcg gc (SEQ ID No: 123)
14	g cgt gga gaC gtc tac ac (SEQ ID No: 124)
15	t cgg aat gGc cag gag g (SEQ ID No: 125)
16	g ctg cct gAc gcc gag (SEQ ID No: 126)
17	c gac gtg gAg gtg tac c (SEQ ID No: 127)
18	g ccg cct gAc gcc gag (SEQ ID No: 128)
19	g cat gga gaC gtc tac ac (SEQ ID No: 129)
20	gc ctc cag aAc ccc atc a (SEQ ID No: 130)
21	g gag tac gcA cgc ttc ga (SEQ ID No: 131)
22	ccg cct gCc gcc gag (SEQ ID No: 132)
23	gg gcg gag Ttg gac acg (SEQ ID No: 133)
24	ac ccc atc aTc gtg gag t (SEQ ID No: 134)
25	gc gtg cgt cTt gtg acc a (SEQ ID No: 135)
26	g ctg ggg cGg cct gac (SEQ ID No: 136)
27	c ctg gag Agg acc cgg (SEQ ID No: 137)

Table 18B-2

Probe No.	Base Sequence
28	aac ggg acC gag cgc g ( SEQ ID No: 138)
29	ag ttg gac aCg gtg tgc a ( SEQ ID No: 139)
30	g gac acc gtA tgc aga ca ( SEQ ID No: 140)
31	g tat cgg gTg gtg acg c ( SEQ ID No: 141)
32	cc cag cat gcC g t gtc tac ( SEQ ID No: 142)
33	t ccc cag cAt gga gac g ( SEQ ID No: 143)
34	ag aag gaa gAc ctg gag ag ( SEQ ID No: 144)
35	g acc gag cTc gtg cgg ( SEQ ID No: 145)
36	g ggg cgg cTt gac gcc ( SEQ ID No: 146)
37	c ttc acc aaT ggg acg ga ( SEQ ID No: 147)
38	gcg gtg acC ccg cag g ( SEQ ID No: 148)
39	t gac cag gaA gag aca gc ( SEQ ID No: 149)
40	t gtg acc aga Tac atc tat aa ( SEQ ID No: 150)
41	gc ggg atc Ttg cag agg ( SEQ ID No: 151)
42	t gac tgg acT ttc cag atc ( SEQ ID No: 152)
43	g cgt ctt gtA acc aga cac ( SEQ ID No: 153)
44	tc aaa gtc cAg tgg ttt cg ( SEQ ID No: 154)
45	gtg tac cgC gcg gtg ac ( SEQ ID No: 155)
46	g gag agg Acc cgg gcg ( SEQ ID No: 156)
47	c gag gtg gGg tac cgc ( SEQ ID No: 157)
48	g cgt ctt gtA acc aga tac ( SEQ ID No: 158)
49	t gta acc aga Tac atc tat aac ( SEQ ID No: 159)
50	cgg cct gtC gcc gag t ( SEQ ID No: 160)
51	c cgg gcg gAg ttg gac ( SEQ ID No: 161)
52	g gtg gcg tTc cgc ggg ( SEQ ID No: 162)
53	gat gcc gag Aac tgg aac ( SEQ ID No: 163)
54	acg ccg cTg ggg cgg ( SEQ ID No: 164)

Table 19A

Allele Number	Probe Number for Detection		
DQA1*010101	0		
DQA1*010102	1		
DQA1*010201	2		
DQA1*010202	3	2	
DQA1*0103	4		
DQA1*010401	5		
DQA1*010402	6	7	
DQA1*0105	8		
DQA1*0106	9		
DQA1*0201	10		
DQA1*030101	11		
DQA1*0302	12		
DQA1*0303	13		
DQA1*040101	14	15	
DQA1*040102	16		
DQA1*050101	17	18	
DQA1*050102	19	20	
DQA1*0502	21		
DQA1*0503	22		
DQA1*0504	23		
DQA1*0505	24		
DQA1*060101	25	26	15
DQA1*060102	27		

Table 19B-1

Allele Number		Probe Number for Detection					
DQB1*050101	0	1	2	3			
DQB1*050102	4						
DQB1*050201	5						
DQB1*050202	6	7	8	4			
DQB1*050301	9	10	4				
DQB1*050302	6	11					
DQB1*0504	7	12					
DQB1*0201	13	14					
DQB1*0202	15	14					
DQB1*0203	16	15					
DQB1*030101	17	18	19	20			
DQB1*030102	17	18					
DQB1*0302	21	22	23	24			
DQB1*030302	18	23	24				
DQB1*030303	25	6	26	18	27	23	
DQB1*0304	17	22	19	20			
DQB1*030501	28	23					
DQB1*030502	6	22	27	29			
DQB1*0306	26	30					
DQB1*0307	31						
DQB1*0308	21	6	22	29			
DQB1*0309	32						
DQB1*0310	6	18	33	19	20		
DQB1*0311	21	6	22	27	29		
DQB1*0312	25	21	6	18	27	23	
DQB1*0313	34						
DQB1*0401	35						
DQB1*0402	36						
DQB1*060101	37						
DQB1*060102	38						
DQB1*060103	39						
DQB1*0602	40	41	42				
DQB1*0603	43	41	42				

Table 19B-2

Allele Number	Probe Number for Detection					
DQB1*060401	27	44				
DQB1*060402	43	45	27	46	47	
DQB1*060501	48	49	27	46	47	
DQB1*060502	48	50	27	46	51	
DQB1*0606	48	49	27	46		
DQB1*0607	43	11	27	46	47	
DQB1*0608	43	45	52			
DQB1*0609	49	27	44			
DQB1*0610	7	41				
DQB1*061101	40	45	11	52		
DQB1*061102	48	49	45	11	41	
DQB1*0612	49	44				
DQB1*0613	40	45	52			
DQB1*0614	43	45	11	41		
DQB1*0615	40	11	27	46	47	
DQB1*0616	53					
DQB1*0617	43	29				
DQB1*0618	48	27	41			
DQB1*0619	25	6	54	11	23	41
DQB1*0620	40	45	11			

Table 20A

Allele Number	Probe Number for Detection		
DQA1*010101	0		
DQA1*010102	1		
DQA1*010201	2		
DQA1*010202	3	2	
DQA1*0103	4		
DQA1*010401	5		
DQA1*010402	6	7	
DQA1*0105	6		
DQA1*0106	8		
DQA1*0201	9		
DQA1*030101	10		
DQA1*0302	11		
DQA1*0303	12		
DQA1*040101	13	14	
DQA1*040102	15		
DQA1*050101	16	17	
DQA1*050102	18	19	
DQA1*0502	20		
DQA1*0503	21		
DQA1*0504	22		
DQA1*0505	23		
DQA1*060101	24	25	14
DQA1*060102	26		

Table 20B-1

Allele Number		Probe Number for Detection					
DQB1*050101	0	1	2	3			
DQB1*050102	4						
DQB1*050201	5						
DQB1*050202	6	7	8	4			
DQB1*050301	9	10	4				
DQB1*050302	6	11					
DQB1*0504	7	12					
DQB1*0201	13	14					
DQB1*0202	15	14					
DQB1*0203	16	15					
DQB1*030101	17	18	19	20			
DQB1*030102	17	18					
DQB1*0302	21	22	23	24			
DQB1*030302	18	23	24				
DQB1*030303	25	6	26	18	27	23	
DQB1*0304	17	22	19	20			
DQB1*030501	28	23					
DQB1*030502	6	22	27	29			
DQB1*0306	26	30					
DQB1*0307	31						
DQB1*0308	21	6	22	29			
DQB1*0309	32						
DQB1*0310	6	18	33	19	20		
DQB1*0311	21	6	22	27	29		
DQB1*0312	25	21	6	18	27	23	
DQB1*0313	34						
DQB1*0401	35						
DQB1*0402	36						
DQB1*060101	37						
DQB1*060102	38						
DQB1*060103	39						
DQB1*0602	40	41	42				
DQB1*0603	43	41	42				



Table 20B-2

Allele Number	Probe Number for Detection					
DQB1*060401	27	44				
DQB1*060402	43	45	27	46	47	
DQB1*060501	48	49	27	46	47	
DQB1*060502	48	50	27	46	51	
DQB1*0606	48	49	27	46		
DQB1*0607	43	11	27	46	47	
DQB1*0608	43	45	52			
DQB1*0609	49	27	44			
DQB1*0610	7	41				
DQB1*061101	40	45	11	52		
DQB1*061102	48	49	45	11	41	
DQB1*0612	49	44				
DQB1*0613	40	45	52			
DQB1*0614	43	45	11	41		
DQB1*0615	40	11	27	46	47	
DQB1*0616	53					
DQB1*0617	43	29				
DQB1*0618	48	27	41			
DQB1*0619	25	6	54	11	23	41
DQB1*0620	40	45	11			

(Example 11)

Probes for identification of HLA-DR allele

Extraction of DNA from 1 ml of human blood was performed using GFX Genomic Blood DNA Purification  
5 Kit from Amersham Biosciences in the same manner as in Example 1.

Next, quantitative PCR was carried out in the same manner as in Example 1 except that probes in the probe list 1 in Tables 21-1 and 21-2 were used and 4  
10  $\mu$ l of the mixed primers consisting of 1  $\mu$ l each of respective solutions of the following primers (10 pmol/ $\mu$ l) and 4  $\mu$ l of ultra pure water were used:

AGAGTACTCCAAGAAACGTG (SEQ ID NO: 822)  
CCGCTGCACCGTGAAGCT (SEQ ID NO: 823)  
15 TCGCTGCACTGTGAAGCT (SEQ ID NO: 824)  
CCTCTGCACTGTGAAGCT (SEQ ID NO: 825).

Referring to Amp Plot and Dissociation curves on a display of 5700 software, it was found that probes 62, 12, and 152 were amplified. Therefore, it  
20 was identified as DRB1\*040502 and DRB1\*130202 referring to the allele-probe list 1 (Tables 23-1 to 23-13).

(Example 12)

Extraction of DNA from 1 ml of human blood was  
25 performed in the same way as in Example 3. PCR of human HLA-DRB exon 2 was then performed in the same manner as in Example 2 except that 6  $\mu$ l of the mixed

primer consisting of 1  $\mu$ l each of the solutions containing the following sequences at 10 pmol/ $\mu$ l respectively, and 9  $\mu$ l of ultra pure water were used:

CCGGATCCTTCGTGTCCCCACAGCACG (SEQ ID NO: 826)

5 AACCCCGTAGTTGTGTCTGCA (SEQ ID NO: 827)

AGAGTACTCCAAGAAACGTG (SEQ ID NO: 822)

CCGCTGCACCGTGAAGCT (SEQ ID NO: 823)

TCGCTGCACTGTGAAGCT (SEQ ID NO: 824)

CCTCTGCACTGTGAAGCT (SEQ ID NO: 825).

10 At the same time, a DNA microarray was prepared to identify the allele in the specimen described above in the same manner as in Example 2, except that probes in the probe list of Tables 22-1 to 22-7 were used to form the probe spots respectively.

15 Then, hybridization was performed using the above specimen and the prepared DNA microarray in the same manner as in Example 2. The fluorometry measurement was conducted with GenePix4000B (Axon).

As a result it was found that probes 59, 133,  
20 and 134 were amplified. Therefore, it was identified as DRB1\*040502 and DRB1\*130202 referring to the allele-probe list 1 (Tables 24-1 to 24-13).

#### Allele list

25 DRB1\*010101 :

atggctgctgaagctccctggaggctcctgcatgacagcgctgacagtacactgatggctgagctccccac  
tggctttggctggggacacccgaccacgtttcttggcagcttaagtgtgaatgtcatttttcaatgggacgga

gcgggtgcggttgc tggAagaTgcatctataaccaagaggagtCcgtgcgcttcgacagcgacgtgggggagtac  
cgggcggtgacggagctggggcgccctgaTgcCgagtactggaacagccagaaggacctctggagcagaggcggg  
ccgcggtggacacctactgcagacacaactacggggttgGtgagagcttcacagtgcagcggcgag (SEQ ID  
NO: 1) ;

5 DRB1\*010102 :

cacgtttcttgtggcagcttaagtttgaatgtcattttcttcaatgggacggagcgggtgcggttgc tggaaagatg  
catctataaccaagaggaAtccgtgcgcttcgacagcgacgtgggggagtagccgggcggtgacggagctggggcgg  
cctgaigccgagtagtggaaacagccagaaggacctctggagcagaggcgggcccgcggtggacacctactgcagac  
acaactacggggttggtagagcttcacagtgcagcggcgag (SEQ ID NO: 2) ;

10 DRB1\*010201 :

ggggacacccgaccacgtttcttgtggcagcttaagtttgaatgtcattttcttcaatgggacggagcgggtgcggt  
tgc tggaaagatgcatctataaccaagaggagtagcgtgcgcttcgacagcgacgtgggggagtagccgggcggtgac  
ggagctggggcgccctgaigccgagtagtggaaacagccagaaggacctctggagcagaggcgggcccgcggtggac  
acctatgcagacacaactacggggCgtggagagcttcacagtgcagcggcgag (SEQ ID NO: 3) ;

15 DRB1\*010202 :

cacgtttcttgtggcagcttaagtttgaatgtcattttcttcaatgggacggagcgggtgcggttgc tggaaagatg  
catctataaccaagaggagtagcgtgcgcttcgacagcgacgtgggggagtagccgggcggtgacggagctggggcgg  
cctgaigccgagtagtggaaacagccagaaggacctctggagcagaggcgggcccgcCgtggacacctatgcagac  
acaactacggggctgtgg (SEQ ID NO: 4) ;

20 DRB1\*0103 :

atggtgtgtctgaagctccctggaggctcctgcatgacagcgctgacagtgcacatgatggtgtctgagctccccac  
tggctttggctggggacacccgaccacgtttcttgtggcagcttaagtttgaatgtcattttcttcaatgggacgga  
gcgggtgcggttgc tggaaagatgcatctataaccaagaggagtagcgtgcgcttcgacagcgacgtgggggagtag  
cgggcggtgacggagctggggcgccctgaigccgagtagtggaaacagccagaaggacAtcctggaagacGAgcggg

25 ccgcggtggacacctactgcagacacaactacggggttggtagagcttcacagtgcagcggcgag (SEQ ID  
NO: 5) ;

DRB1\*0104 :

ggggacacccgaccacgtttcttggcagcttaagtttgaatgtcattttcttcaatgggacggagcgggtgcggt  
tgc tggaaagatgcatctataaccaagaggagtcggtgcgcttcgacagcgacgtgggggagtagcgggcggtgac  
ggagctggggcggcctgatgccgagtagtgaacagccagaaggacctcctggagcagaggcgggccggtggac  
aaTtactgcagacacaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 6) ;

5 DRB1\*0105 :

cacgtttcttggcagcttaagtttgaatgtcattttcttcaatgggacggagcgggtgcggttgc tggaaagatg  
catctataaccaagaggagtcggtgcgcttcgacagcgacgtAgggagtagcgggcggtgacggagctggggcgg  
cctgatgccgagtagtgaacagccagaaggacctcctggagcagaggcgggccggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 7) ;

10 DRB1\*0106 :

cacgtttcttggcagcttaagtttgaatgtcattttcttcaatgggacggagcgggtgcggttgc tggaaagatg  
catctataaccaagaggagtcggtgcgcttcgacagcgacgtgggggagtagcgggcggtgacggagctggggcgg  
cctgatgccgagtagtgaacagccagaaggacctcctggagcaggCgcgggccggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 8) ;

15 DRB1\*0107 :

cacgtttcttggGagcttaagtttgaatgtcattttcttcaatgggacggagcgggtgcggttgc tggaaagatg  
catctataaccaagaggagtcggtgcgcttcgacagcgacgtgggggagtagcgggcggtgacggagctggggcgg  
cctgatgccgagtagtgaacagccagaaggacctcctggagcagaggcgggccggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 9) ;

20 DRB1\*0108 :

cacgtttcttggcagcttaagtttgaatgtcattttcttcaatgggacggagcgggtgcggttgc tggaaagatg  
catctataaccaagaggagtagtgcgcttcgacagcgacgtgggggagtagcgggcggtgacggagctggggcgg  
cctgatgccgagtagtgaacagccagaaggacctcctggagcagaggcgggccggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 10) ;

25 DRB1\*0109 :

cacgtttcttggcagcttaagtttgaatgtcattttcttcaatgggacggagcgggtgcggttgc tggaaagatg  
catctataaccaagaggagtcggtgcgcttcgacagcgacgtgggggagtagcgggcggtgacggagctggggcgg

cctgatgccgagtlactggaacagccagaaggacctcctggagcagGCgcggggccgcggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 1 1) ;

DRB1\*0110 :

cacgtttcttggcagcttaagtttgaatgtcatttcttcaatgggacggagcgggtgcggttgctggaaagatg  
5 calclataaccaagaggagtcctgtgcgttcgacagcgacgtgggggagtagccggcggtgacggagctggggcgg  
cctgatgccgagtlactggaacagccagaaggacctcctggagcagaAgcggggccgcggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 1 2) ;

DRB1\*030101 :

ggggacaccagaccacgtttcttggagtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggt  
10 AccitggacagatacttcCataaccaggaggagAACgtgcgcttcgacagcgacgtgggggagtagccggcggtgac  
ggagctggggcggcctgatgccgagtlactggaacagccagaaggacctcctggagcagaagcggggccGggtggac  
aActactgcagacacaactacggggttggtGgagagcttcacagtcagcggcgag (SEQ ID NO: 1 3) ;

DRB1\*030102 :

cacgtttcttggagtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggtacctggacagata  
15 ctltccataaccaggaggagaacgtgcgcttcgacagcgacgtgggggagtagccggcggtgacggagctggggcgg  
cctgatgccgagtlactggaacagccagaaggacctcctggagcagaagcggggccGggtggacaaTtactgcagac  
acaactacggggttggtGgagagcttcacagtcagcg (SEQ ID NO: 1 4) ;

DRB1\*030201 :

ggggacaccagaccacgtttcttggAgtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggt  
20 tcctggaGagatacttcCataaccaggaggagAACgtgcgcttcgacagcgacgtgggggagtagccggcggtgac  
ggagctggggcggcctgatgccgagtlactggaacagccagaaggacctcctggagcagaagcggggccGggtggac  
aActactgcagacacaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 1 5) ;

DRB1\*030202 :

ggggacaccagaccacgtttcttggagtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggt  
25 tcctggaGagatacttcCataaccaggaggagAACgtgcgcttcgacagcgacgtgggggagtagccggcggtgac  
ggagctggggcggcctgatgccgagtlactggaacagccagaaggacctcctggagcagaagcggggccGggtggac  
aaTtactgcagacacaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 1 6) ;

DRB1\*0303 :

tactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttccctggaGagatacttcCataaccagg  
aggagAACgtgcgcttcgacagcgacgtgggggagtlaccgggcggtagcggagctggggcgccctgatgccgagta  
ctggaacagccagaaggacctccctggagcagaagcggggccGggtggacaActactgcagacacaactacgggggt  
5 gtGgagagcttcacagtgcagcggcga (SEQ ID NO: 17) ;

DRB1\*0304 :

cacgttcttggagtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggtAccctggacagata  
cttcCataaccaGgaggagtcggtgcgcttcgacagcgacgtgggggagTccgggcggtagcggagctggggcgg  
ccctgatgccgagtagctggaacagccagaaggacctccctggagcagaagcggggccGggtggacaActactgcagac  
10 acaactacgggggtgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 18) ;

DRB1\*030501 :

cacgttcttggagtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggtAccctggacagata  
cttcCataaccaggaggagAACgtgcgcttcgacagcgacgtgggggagTccgggcggtagcggagctggggcgg  
ccctgatgccgagtagctggaacagccagaaggacctccctggagcagaagcggggccGggtggacaActactgcagac  
15 acaactacgggggtggtagagcttcacagtgcagcggcgag (SEQ ID NO: 19) ;

DRB1\*030502 :

cacgttcttggagtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggtacctggacagata  
cttcCataaccaggaggagaacgtgcgcttcgacagcgacgtgggggagTccgggcggtagcggagctggggcgg  
ccctgatgccgagtagctggaacagccagaaggacctccctggagcagaagcggggccgggtggacaActactgcagac  
20 acaactacgggggtggtagagcttcacGgtgcagcggcgag (SEQ ID NO: 20) ;

DRB1\*0306 :

tcttggagtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggtAccctggaCagatacttcC  
ataaccaggaggagAACgtgcgcttcgacagcgacgtgggggagtlaccgggcggtagcggagctggggcgccctga  
tgccgagtagctggaacagccagaaggacctccctggagcagaagcggggccGggtggacaActactgcagacacaac  
25 tacgggggtgtGgagagcttcacagtgcag (SEQ ID NO: 21) ;

DRB1\*0307 :

ggggacaccagaccacgttcttggagtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggt

tcctggacagatacttcCataaccaggaggagAACgtgcgcttcgacagcgacgtgggggagTccgggcggtagac  
ggagctggggcggcctgatgccgagtagtggaaacagccagaaggacctccctggagcagaagcggggccGggtggac  
aActactgcagacacaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 2) ;

DRB1\*0308 :

5 ggggacaccagaccacgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggt  
acctggacagatacttccataaccaggaggagAACgtgcgcttcgacagcgacgtgggggagTccgggcggtagac  
ggagctggggcggcctgatAGgagtagtggaaacagccagaaggacctccctggagcagaagcggggccGggtggac  
aActactgcagacacaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 3) ;

DRB1\*0309 :

10 ttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggtacctggacagatacttccata  
accGggaggagaacgtgcgcttcgacagcgacgtgggggagTccgggcggtagcggagctggggcggcctgatgc  
cgagtactggaaacagccagaaggacctccctggagcagaagcggggccgggtggacaactactgcagacacaactac  
ggggttggtagagcttcacagtcagcgg (SEQ ID NO: 2 4) ;

DRB1\*0310 :

15 ggggacaccagaccacgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggt  
acctggacagatacttccataaccaggaggagaacgtgcgcttcgacagcgacgtgggggagTccgggcggtagac  
ggagctggggcggcctgCtgcggagcac tggaaacagccagaaggacctccctggagcagaagcggggccGggtggac  
aActactgcagacacaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 5) ;

DRB1\*0311 :

20 cacgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggtAccctggacagata  
cttccataaccaggaggagaacgtgcgcttcgacagcgacgtgggggagTccgggcggtagcggagctggggcgg  
cctgatgccgagtagtggaaacagccagaaggacctccctggagcagaagcggggcCAGgtggacaActactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcga (SEQ ID NO: 2 6) ;

DRB1\*0312 :

25 ttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggtacctggacagatacttccata  
accaggaggagaacgtgcgcttcgacagcgacgtgggggagTccgggcggtagcggagctggggcggcctagCgc  
cgagtactggaaacagccagaaggacctccctggagcagaagcggggccGggtggacaActactgcagacacaactac



ggggltgtGgag(SEQ ID NO: 2 7) ;

DRB1\*0313 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggtacctggacagata  
cttccataaccaggaggagaacgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
5 cctgatgccgagtcctggaacagccagaaggacctcctggagcagaagcggggccGggtggacaActactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag(SEQ ID NO: 2 8) ;

DRB1\*0314 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggtAcctggacagata  
cttcCataaccaggaggagAAcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
10 cctgatgccgagtcctggaacagccagaaggacctcctggagcagaagcggggccGggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag(SEQ ID NO: 2 9) ;

DRB1\*0315 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggtAcctggacagata  
cttcCataaccaggaggagAAcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
15 cctgatgccgagtcctggaacagccagaaggacctcctggagcagaagcggggccGggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcga(SEQ ID NO: 3 0) ;

DRB1\*0316 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggtacctggacagata  
cttccataaccaggaggagaacgtgcgcttcgacagcgacgtgggggagttcTgggcggtgacggagctggggcgg  
20 cctgatgccgagtcctggaacagccagaaggacctcctggagcagaagcggggccgggtggacaactactgcagac  
acaactacggggttgtg(SEQ ID NO: 3 1) ;

DRB1\*0317 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttcctggaCagata  
cttctataaccaagaggagAcgtgcgcttcgacagcgacgtgggggagttaccgggcggtgaGggagctggggcgg  
25 cctgatgccgagtcctggaacagccagaaggacctcctggagcagaagcggggccagggtggacaaTtactgcagac  
acaactacggggttgtgagagcttcacagtcagcggcgag(SEQ ID NO: 3 2) ;

DRB1\*0318 :

cacgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggtacctggacagata  
cttccataaaccaggaggagaacgtgcgcttcgacagcgacgtgCgggagttccgggcggtgacggagctggggcgg  
cctgatgccgagttactggaacagccagaaggacctcctggagcagaagcggggccgggtggacaactactgcagac  
acaactacgggggtgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 3 3) ;

5 DRB1\*0319 :

cacgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggtacctggacagata  
cttccataaaccaggaggagaacgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
cctgatgccgagttactggaacagccagaaggacAtcctggagcagaagcggggccGggtggacaActactgcagac  
acaactacgggggtgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 3 4) ;

10 DRB1\*0320 :

cacgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggtacctggacagata  
cttccataaaccaggaggagaacgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
cctgatgccgagttactggaacagccagaaggacctcctggagcagaagcggggccgggtggacaActactgcagac  
acaactacggggCtgtggagagcttcacagtcagcgg (SEQ ID NO: 3 5) ;

15 DRB1\*0321 :

cgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggtacctggacagatact  
tccataaaccaggaggagttcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcc  
tgatgccgagttactggaacagccagaaggacctcctggagcagaagcggggccGggtggacaActactgcagacac  
aactacgggggtgtGgagagcttcacagtcagcggcga (SEQ ID NO: 3 6) ;

20 DRB1\*0322 :

tttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggtacctggacagatacttc  
Gataaaccaggaggagaacgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctg  
atgccgagttactggaacagccagaaggacctcctggagcagaagcggggccgggtggacaactactgcagacacaa  
ctacgggggtgtggagagcttcacagtcagcggcgag (SEQ ID NO: 3 7) ;

25 DRB1\*0323 :

cacgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggtacctggacagata  
cttccataaaccGggaggagaacgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg

cctgatgccgagtlactggaacagccagaaggacctcctggagcagaagcggggccgggtggacaactactgcagac  
acaactacgggggtgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 38) ;

DRBI\*0324 :

cacgtttcttggagtlactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttcttggacagata  
5 ctttcataaccaggaggagaacgtgcgcttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcgg  
cctgatgccgagtlactggaacagccagaaggacctcctggagcagaagcggggcCAGgtggacaaTtactgcagac  
acaactacgggggtgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 39) ;

DRBI\*0325 :

cacgtttcttggagtlactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggtAcctggacagata  
10 ctttcCataaccaGaggagTAcgtgcgcttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcgg  
cctgatgccgagtlactggaacagccagaaggacctcctggagcagaagcggggcGggtggacaActactgcagac  
acaactacgggggtgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 40) ;

DRBI\*040101 :

atgggtgtgtctgaagTtccctggaggctcctgcatggcagctctgacagtgcacactgatgggtgtgagctccccac  
15 tggctttggctggggacacccgaccacgtttcttggagcagggttaacatgagtgatcttcttcaacgggacgga  
gcgggtgcggttcttggacagatacttctatcaccaagaggagtlacgtgcgcttcgacagcgacgtgggggagtlac  
cgggcggtgacggagctggggcggcctgatgccgagtlactggaacagccagaaggacctcctggagcagaAgcggg  
ccgcgggtggacacctactgcagacacaactacgggggtgggtgagagcttcacagtgcagcggcgag (SEQ ID  
NO: 41) ;

20 DRBI\*040102 :

cacgtttcttggagcagggttaacatgagtgatcttcttcaacgggacggagcgggtgcggttcttggacagata  
cttctatcaccaagaAgagtlacgtgcgcttcgacagcgacgtgggggagtaccgggcggtgacggagctggggcgg  
cctgatgccgagtlactggaacagccagaaggacctcctggagcagaagcggggcgggtggacacctactgcagac  
acaactacgggggtgggtgagagcttcacagtgcagcggcgag (SEQ ID NO: 42) ;

25 DRBI\*0402 :

atgggtgtgtctgaagTtccctggaggctcctgcatggcagctctgacagtgcacactgatgggtgtgagctccccac  
tggctttggctggggacacccgaccacgtttcttggagcagggttaacatgagtgatcttcttcaacgggacgga

gcgggtgcggttccctggacagatacttctatcaccaagaggagtacgtgcgcttcgacagcgacgtgggggagttac  
cgggcgggtgacggagctggggcggccctgatgccgagttactggaacagccagaaggacattccctggaagacgAgcggg  
ccgcggtggacacctactgcagacacaactacggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID  
NO: 4 3) ;

5 DRB1\*040301 :

ggggacacccgaccacgtttcttggagcaggttaaaCatgagtgctcatttcttcaacgggacggagcgggtgcggt  
tccctggacagatacttctatcaccaagaggagtAcgtgcgcttcgacagcgacgtgggggagttaccgggcgggtgac  
ggagctggggcggccctgatgccgagttactggaacagccagaaggacctccctggagcagaggcgggcccAggtggac  
acctactgcagacacaactacggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 4 4) ;

10 DRB1\*040302 :

cacgtttcttggagcaggttaaaCatgagtgctcatttcttcaacgggacggagcgggtgcggttccctggacagata  
cttctatcaccaagaggagtacgtgcgcttcgacagcgacgtgggggagttaccgggcgggtgacggagctggggcgg  
cctgacgcTgagttactggaacagccagaaggacctccctggagcagaggcgggcccAggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 4 5) ;

15 DRB1\*0404 :

atgggtgtgtctgaagTtcccggaggctccctgcatggcagcttcgacagtacactgatgggtgtgagctccccac  
tggctttggctggggacacccgaccacgtttcttggagcaggttaaacatgagtgctcatttcttcaacgggacgga  
gcgggtgcggttccctggacagatacttctatcaccaagaggagtacgtgcgcttcgacagcgacgtgggggagttac  
cgggcgggtgacggagctggggcggccctgatgccgagttactggaacagccagaaggacctccctggagcagaggcggg

20 ccgcggtggacacctactgcagacacaactacggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID  
NO: 4 6) ;

DRB1\*040501 :

ggggacacccgaccacgtttcttggagcaggttaaaCAtgagtgctcatttcttcaacgggacggagcgggtgcggt  
tccctggacagatacttctatCaccaagaggagtAcgtgcgcttcgacagcgacgtgggggagttaccgggcgggtgac

25 ggagctggggcggccctAGcggcagttactggaacagccagaaggacctccctggagcagaggcgggcccgggtggac  
acctactgcagacacaactacggggttggtgagagcttcacagtgcagcggcga (SEQ ID NO: 4 7) ;

DRB1\*040502 :

cacgtttcttggagcaggtaaacaatgagtgatcatttcttcaacgggacggagcgggtgcggttccitggacagata  
cttctatcaccaagaggagtagctgcgttcgacagcgacgtgggggagtagcgggcggtagcggagctggggcgg  
cctagcgccgagtagtggaacagccagaaggaccttctggagcagaggcgggccggtggacacctactgcagac  
acaactacgggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 48) ;

5 DRB1\*040503 :

cacgtttcttggagcaggtaaacaatgagtgatcatttcttcaacgggacggagcgggtgcggttccitggacagata  
cttctatcaccaagaggagtagctgcgttcgacagcgacgtgggggagtagcgggcggtagcggagctggggcgg  
cctagcgccgagtagtggaacagccagaaggaccttctggagcagaggcgggccggtggacacctactgcagac  
acaactacgggggttggtgagagcttcacagtcagcgcgag (SEQ ID NO: 49) ;

10 DRB1\*040504 :

cacgtttcttggagcaggtaaacaatgagtgatcatttcttcaacgggacggagcgggtgcggttccitggacagata  
cttctatCaccaagaggagtagctgcgttcgacagcgacgtgggggagtagcgggcggtagcggagctggggcgg  
cctagCgccgagtagtggaacagccagaaggaccttctggagcagaggcgggccggtggacacctactgcagac  
acaactacgggggttggtgagagcttcacGgtgcagcggcgag (SEQ ID NO: 50) ;

15 DRB1\*0406 :

ggggacacccgaccacgtttcttggagcaggtaaacaatgagtgatcatttcttcaacgggacggagcgggtgcggt  
tccitggacagatacttctatCaccaagaggagtagctgcgttcgacagcgacgtgggggagtagcgggcggtagc  
ggagctggggcggccctgatgccgagtagtggaacagccagaaggaccttctggagcagaggcgggccgAggtggac  
acctactgcagacacaactacgggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 51) ;

20 DRB1\*040701 :

ggggacacccgaccacgtttcttggagcaggtaaacaatgagtgatcatttcttcaacgggacggagcgggtgcggt  
tccitggacagatacttctatcaccaagaggagtagctgcgttcgacagcgacgtgggggagtagcgggcggtagc  
ggagctggggcggccctgatgccgagtagtggaacagccagaaggaccttctggagcagaggcgggccgAggtggac  
acctactgcagacacaactacgggggttggtgagagcttcacagtcagcggcga (SEQ ID NO: 52) ;

25 DRB1\*040702 :

cacgtttcttggagcaggtaaacaatgagtgatcatttcttcaacgggacggagcgggtgcggttccitggacagata  
cttctatcaccaagaggagtagctgcgttcgacagcgacgtgggggagtagcgggcggtagcggagctggggcgg

ccatgatgccgagttacttggaacagccagaaggacctccctggagcagagAcggggccgaggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtcagcgg(SEQ ID NO: 53) ;

DRBI\*0408 :

ttctctggagcaggttaaACAgtggtgcatcttcttcaacgggacggagcgggtgcggttccctggacagatacttc  
5 tatCaccaagaggagtAcgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcggcctg  
atgccgagttacttggaacagccagaaggacctccctggagcagaggcgggccggtggacacctactgcagacacaa  
ctacggggttggtgagagcttcacagtcagcggcgag(SEQ ID NO: 54) ;

DRBI\*0409 :

tgagtggtcatcttcttcaacgggacggagcgggtgcggttccctggacagatacttctatCaccaagaggagtacgtg  
10 cgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcggcctaGgccgagttacttggaacagcc  
agaaggacctccctggagcagaAgcgggccggtggacacctactgcagacacaaactacggggttggtgagag(SE  
Q ID NO: 55) ;

DRBI\*0410 :

ttctctggagcaggttaaacaatgagtggtcatcttcttcaacgggacggagcgggtgcggttccctggacagatacttc  
15 tatCaccaagaggagtAcgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcggccta  
Ggccgagttacttggaacagccagaaggacctccctggagcagaggcgggccggtggacacctactgcagacacaa  
ctacggggttgtGgagagcttcacagtcagcggcgag(SEQ ID NO: 56) ;

DRBI\*0411 :

atgggtgtgtctgaagTtccctggaggctccctgcatggcagctctgacagtgacactgatgggtgtgagctccccac  
20 tggctttggctggggacacccgaccagttcttggagcaggttaaacaatgagtggtcatcttcttcaacgggacgga  
gcgggtgcggttccctggacagatacttctatcaccaagaggagtlacgtgcgcttcgacagcgacgtgggggagtlac  
cgggcggtgacggagctggggcggcctagcggcgagttacttggaacagccagaaggacctccctggagcagaggcggg  
ccgAggtggacacctactgcagacacaaactacggggttgtGgagagcttcacagtcagcggcgag(SEQ ID  
NO: 57) ;

25 DRBI\*0412 :

ttctctggagcaggttaaacaatgagtggtcatcttcttcaacgggacggagcgggtgcggttccctggacagatacttct  
atCaccaagaggagtlacgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcggcctaG

cgccgagttacttggaacagccagaaggacAtcctggaagacaggcgggccTggtggacacctactgcagacacaac  
tacggggttgtGgagagcttcacagtcagcgg(SEQ ID NO: 5 8) ;

DRB1\*0413 :

catgagtgctatttcttcaacgggacggagcgggtgcggttccctggacagatacttctatCaccaagaggagtacg  
5 tgcgcttcgacagcgacgtgggggagttaccgggcggtgacggagctggggcgccctgatgccgagttacttggaacag  
ccagaaggaccttctggagcagaAgcgggcccgcggtggacacctactgcagacacaactacggggttgtGgagagc  
ttcaca(SEQ ID NO: 5 9) ;

DRB1\*0414 :

tgagtgctatttcttcaacgggacggagcgggtgcggttccctggacagatacttctatCaccaagaggagtacgtg  
10 cgcttcgacagcgacgtgggggagttaccgggcggtgacggagctggggcgccctgatgccgagttacttggaacagcc  
agaaggacAtcctggaagacGAgcgggcccgcggtggacacctactgcagacacaactacggggttggtagag(SE  
Q ID NO: 6 0) ;

DRB1\*0415 :

cacgtttcttggagcaggttaaaCatgagtgctatttcttcaacgggacggagcgggtgcggttccctggacagata  
15 ctctatcaccaagaggagtacgtgcgcttcgacagcgacgtgggggagttaccgggcggtgacggagctggggcg  
cctgatgaGgagttacttggaacagccagaaggacTtccctggaagaCaggcgggcccgcggtggacacctactgcagac  
acaactacggggttgtGgagag(SEQ ID NO: 6 1) ;

DRB1\*0416 :

atgagtgctatttcttcaacgggacggagcgggtgcggttccctggacagatacttctatCaccaagaggagtacgt  
20 gcgcttcgacagcgacgtgggggagttaccgggcggtgacggagctggggcgccctgatgccCagttacttggaacagc  
cagaaggaccttctggagcagaagcgggcccgcggtggacacctactgcagacacaactacggggttggtg(SEQ  
ID NO: 6 2) ;

DRB1\*0417 :

atgagtgctatttcttcaacgggacggagcgggtgcggttccctggacagatacttctatCaccaagaggagtacgt  
25 gcgcttcgacagcgacgtgggggagttaccgggcggtgacggagctggggcgccctGcgccgagttacttggaacagc  
cagaaggaccttctggagcagaggcgggcccAggtggacacctactgcagacacaactacggggttggt(SEQ ID  
NO: 6 3) ;

DRB1\*0418 :

atgagtgatcatttcttcaacgggacggagcgggtgcggttccctggacagatacttctatCaccaagaggagtagct  
gcgcttcgacagcgacgtgggggagtagcgggcggtagcggagctggggcggccctgatgccgagtagtggaaacagc  
cagaaggacAtcctgggaagacaggcgggcccTggtggacacctactgcagacacaactacgggggtgtGgagagct  
5 tccagtgca (SEQ ID NO: 64) ;

DRB1\*0419 :

tttcttggagcagggttaaACAtgagtgatcatttcttcaacgggacggagcgggtgcggttccctggacagatacttct  
tatCaccaagaggagtagcgtgcgcttcgacagcgacgtgggggagtagcgggcggtagcggagctggggcggccctg  
atgccgagtagtggaaacagccagaaggacctcctggagcagaggcgggcccgcggtggacacctactgcagacacaa  
10 ctacgggggttggtgagagcttcacagtgacggcgag (SEQ ID NO: 65) ;

DRB1\*0420 :

atgagtgatcatttcttcaacgggacggagcgggtgcggttccctggacagatacttctatCaccaagaggagtagcgt  
gcgcttcgacagcgacgtgggggagtagcgggcggtagcggagctggggcggccctgatgccgagtagtggaaacagc  
cagaaggacctcctggagcagaggcgggcccAggtggacacctactgcagacacaactacgggggttggtg (SEQ  
15 ID NO: 66) ;

DRB1\*0421 :

gagcagggttaaacaatgagtgatcatttcttcaacgggacggagcgggtgcggttccctggacagatacttctatCacc  
aagaggagtagcgtgcgcttcgacagcgacgtgggggagtagcgggcggtagcggagctggggcggccctgatgccga  
gtactggaaacagccagaaggacctcctggagcagaAgcgggcccgcggtggacacctactgcagacacaactacggg  
20 gtgtggtgagagcttcacagt (SEQ ID NO: 67) ;

DRB1\*0422 :

gagcagggttaaacaatgagtgatcatttcttcaacgggacggagcgggtgcggttccctggacagatacttctatCacc  
aagaggagtagcgtgcgcttcgacagcgacgtgggggagtagcgggcggtagcggagctggggcggccctgatgccga  
gtactggaaacagccagaaggacctcctggagcagaagcgggcccGggtggacaActactgcagacacaactacggg  
25 gtgtGgagagcttcaca (SEQ ID NO: 68) ;

DRB1\*0423 :

cacgtttcttggagcagggttaaacaatgagtgatcatttcttcaacgggacggagcgggtgcggttccctggacagata



cttctatcaccaagaggagtagtgcgcttcgacagcgacgtgggggagtagcggcggtgacggagctggggcgg  
ccgatgccgagtagtggaacagccagaaggacctcctggagcagaggcgggccgggtggacacctactgcagac  
acaactacggggttggtgagagattcacagtagcagggcgag (SEQ ID NO: 69) ;

DRB1\*0424 :

5 cactttcttggagcagggttaaacaatgagtgatcttcttcaacgggacggagcgggtgcggttccctggacagata  
cttctatcaccaagaggagtagtgcgcttcgacagcgacgtgggggagtagcggcggtgacggagctggggcgg  
cctaGcggcagtagtggaacagccagaaggacctcctggagcGgaggcgggccgggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtagcagggcgag (SEQ ID NO: 70) ;

DRB1\*0425 :

10 ttggagcagggttaaacaatgagtgatcttcttcaacgggacggagcgggtgcggttccctggacagatacttctatC  
accaagaggagtagtgcgcttcgacagcgacgtgggggagtagcggcggtgacggagctggggcggccgatgc  
cgagtagtggaacagccagaaggacTtcttgaagacaggcgggccTggtggacacctactgcagacacaactac  
ggggttggtGgagag (SEQ ID NO: 71) ;

DRB1\*0426 :

15 cactttcttggagcagggttaaacaatgagtgatcttcttcaacgggacggagcgggtgcggttccctggacagata  
cttctatcaccaagaggagtagtgcgcttcgacagcgacgtgggggagtagcggcggtgacggagctggggcgg  
ccgatAccgagtagtggaacagccagaaggacctcctggagcagaagcgggccgggtggacacctactgcagac  
acaactacggggttggtg (SEQ ID NO: 72) ;

DRB1\*0427 :

20 cactttcttggagcagggttaaacaatgagtgatcttcttcaacgggacggagcgggtgcggttccctggacagata  
cttctatcaccaagaggagtagtgcgcttcgacagcgacgtgggggagtagcggcggtgacggagctggggcgg  
ccgatgccgagtagtggaacagccagaaggacctcctggagcagaggcgggccgAggtggacacctactgcagac  
acaactacggggCtgggagagcttcacagtg (SEQ ID NO: 73) ;

DRB1\*0428 :

25 cactttcttggagcagggttaaacaatgagtgatcttcttcaacgggacggagcgggtgcggttccctggacagata  
cttctatCaccaagaggagtagtgcgcttcgacagcgacgtgggggagtagcggcggtgacggagctggggcgg  
cctaGcggcagtagtggaacagccagaaggacctcctggagcagaggcgggccgggtggacacctactgcagac

acaactacgggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 7 4) ;

DRB1\*0429 :

cacgtttcttggagcagggttaaacaatgagtgatcatttcttcaacgggacggagcgggtgcggttccctggacagata  
cttctatcaccaagaggagtagctgcgcttcgacagcgacgtgggggagtagcgggcggtgaTggagctggggcgg

5 cctagcgccgagtagctggaacagccagaaggaccttctggagcagaggcgggcccgggtggacacctactgcagac  
acaactacgggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 7 5) ;

DRB1\*0430 :

cacgtttcttggagcagggttaaacaatgagtgatcatttcttcaacgggacggagcgggtgcggttccctggacagata  
cttctatcaccaagaggagtagctgcgcttcgacagcgacgtgggggagtagcgggTggtgacggagctggggcgg

10 cctagcgccgagtagctggaacagccagaaggaccttctggagcagaggcgggcccgggtggacacctactgcagac  
acaactacgggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 7 6) ;

DRB1\*0431 :

cacgtttcttggagcagggttaaaCatgagtgatcatttcttcaacgggacggagcgggtgcggttccctggacagata  
cttctatCaccaagaggagtagctgcgcttcgacagcgacgtgggggagtagcgggcggtgacggagctggggcgg

15 cctgatgccgagtagctggaacagccagaaggaccttctggagcagaggcgggcccTggtggacacctactgcagac  
acaactacgggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 7 7) ;

DRB1\*0432 :

tggagcagggttaaacaatgagtgatcatttcttcaacgggacggagcgggtgcggttccctggacagatacttctatc  
accaagaggagtagctgcgcttcgacagcgacgtgggggagtagcgggcggtgacggagctggggcgccctgatgc

20 cgagtagctggaacagccagaaggaccttctggagcagaggcAggcccgggtggacacctactgcagacacacacac  
gggggttggag (SEQ ID NO: 7 8) ;

DRB1\*0433 :

cacgtttcttggagcagggttaaacaatgagtgatcatttcttcaacgggacggagcgggtgcggttccctggacagata  
cttctatcaccaagaggagtagctgcActtcgacagcgacgtgggggagtagcgggcggtgacggagctggggcgg

25 cctgatgccgagtagctggaacagccagaaggaccttctggagcagaagcgggcccgggtggacacctactgcagac  
acaactacgggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 7 9) ;

DRB1\*0434 :

t t t c t t g g a g c a g g t t a a a C C t g a g t g t c a t t t c t t c a a c g g g a c g g a g c g g g t g c g g t t c c t g g a c a g a t a c t t c  
t a t c a c c a a g a g g a g t A c g t g c g c t t c g a c a g c g a c g t g g g g a g t a c c g g g c g g t g a c g g a g c t g g g c g g c c t g  
a t g c c g a g t a c t g g a a c a g c c a g a a g g a c c t c c t g g a g c a g a A g c g g g c c g c g g t g g a c a c c t a c t g c a g a c a a  
c t a c g g g g t t g g t g a (SEQ ID NO: 8 0) ;

## 5 DRB1\*0435 :

c a c g t t t c t t g g a g c a g g t t a a a C a t g a g t g t c a t t t c t t c a a c g g g a c g g a g c g g g t g c g g t t c c t g g a c a g a t a  
c t t c t a t c a c c a a g a g g a g t a c g t g c g c t t c g a c a g c g a c g t g g g g a g t T c c g g g c g g t g a c g g a g c t g g g c g g  
c c t g a t g c c g a g t a c t g g a a c a g c c a g a a g g a c c t c c t g g a g c a g a A g c g g g c c g c g g t g g a c a c c t a c t g c a g a c  
a c a a c t a c g g g g t t g g t g a g a g c t t c a c a g t g (SEQ ID NO: 8 1) ;

## 10 DRB1\*0436 :

c a c g t t t c t t g g a g c a g g t t a a a C a t g a g t g t c a t t t c t t c a a c g g g a c g g a g c g g g t g c g g t t c c t g g a c a g a t a  
c t t c t a t c a c c a a g a g g a g t a c g t g c g c t t c g a c a g c g a c g t g g g g a g t a c c g g g c g g t g a c g g a g c t g g g c g g  
c c t g a t g c c g a g t a c t g g a a c a g c c a g a a g g a c t t c c t g g a a g a C a g g c g g g c c g c g g t g g a c a c c t a c t g c a g a c  
a c a a c t a c g g g g t t g t G g a g a g c t t c a c a g t g c a g c g g c g a g (SEQ ID NO: 8 2) ;

## 15 DRB1\*0437 :

c a c g t t t c t t g g a g c a g g t t a a a C a t g a g t g t c a t t t c t t c a a c g g g a c g g a g c g g g t g c g g t t c c t g g a c a g a t a  
c t t c t a t c a c c a a g a g g a g t a c g t g c g c t t c g a c a g c g a c g t g g g g a g t a c c g g g c g g t g a c g g a g c t g g g c g g  
c c t g a t g c c g a g t a c t g g a a c a g c c a g a a g g a c c t c c t g g a a g a C A g c g g g c c g c g g t g g a c a c c t a c t g c a g a c  
a c a a c t a c g g g g t t g t G g a g a g c t t c a c a g t g c a g c g g c g a g (SEQ ID NO: 8 3) ;

## 20 DRB1\*0438 :

c a c g t t t c t t g g a g c a g g t t a a a C a t g a g t g t c a t t t c t t c a a c g g g a c g g a g c g g g t g c g g t t c c t g g a c a g a t a  
c t t c t a t c a c c a a g a g g a g t a c g t g c g c t t c g a c a g c g a c g t g g g g a g t a c c g g g c g g t g a c g g a g c t g g g c g g  
c c t g a t g c c g a g t a c t g g a a c a g c c a g a a g g a c A t c c t g g a g c a g a A g c g g g c c g c g g t g g a c a c c t a c t g c a g a c  
a c a a c t a c g g g g t t g g t g a g a g c t t c a c a g t g c a g c g g c g a g (SEQ ID NO: 8 4) ;

## 25 DRB1\*0439 :

c a c g t t t c t t g g a g c a g g t t a a a C a t g a g t g t c a t t t c t t c a a c g g g a c g g a g c g g g t g c g g t t c c t g g a c a g a t a  
c t t c t a t c a c c a a g a g g a g t a c g t g c g c t t c g a c a g c g a c g t g g g g a C t a c c g g g c g g t g a c g g a g c t g g g c g g

cctgatgccgagtlactiggaacagccagaaggacctccctggagcagaggcgggcccagggtggacacctactgcagac  
acaactacggggttgtggagagcttcacagtcagcgg (SEQ ID NO: 85) ;

DRB1\*0440 :

cacgtttcttggagcagggttaacatgagtgctatttcttcaacgggacggagcgggtgcggttccctggacagata  
5 cttctatcaccaagaggagtagctgcgttcgacagcgacgtgggggagtagcggcggtgacggagctggggcgg  
cctgatgCcgagtlactiggaacagccagaaggacctccctggagcagaggcgggcccgggtggacacctactgcagac  
acaactacggggttgtggagagcttcacagtcagcgg (SEQ ID NO: 86) ;

DRB1\*0441 :

cacgtttcttggagcagggttaaaCatgagtgctatttcttcaacgggacggagcgggtgcggttccctggacagata  
10 cttctatcaccaagaggagAACgtgcgttcgacagcgacgtgggggagtagcggcggtgacggagctggggcgg  
cctgatgccgagtlactiggaacagccagaaggacctccctggagcagaggcgggcccAggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcgg (SEQ ID NO: 87) ;

DRB1\*0442 :

cacgtttcttggagcagggttaaaCatgagtgctatttcttcaacgggacggagcgggtgcggttccctggacagata  
15 cttctatcaccaagaggagtagctgcgttcgacagcgacgtgggggagtagcggcggtgacggagctggggcgg  
cctgatgccgagtlactiggaacagccagaaggacctccctggagcagaggcgggcccgggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 88) ;

DRB1\*0443 ;

cacgtttcttggagcagggttaaaCatgagtgctatttcttcaacgggacggagcgggtgcggttccctggacagata  
20 cttctatCaccaagaggagtagctgcgttcgacagcgacgtgggggagtagcggcggtgacggagctggggcgg  
cctgatgccgagtlactiggaacagccagaaggacctccctggagcagaggcgggcccgggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtcagcgg (SEQ ID NO: 89) ;

DRB1\*0444 :

cacgtttcttggagcagggttaacatgagtgctatttcttcaacgggacggagcgggtgcggttccctggacagata  
25 cttctatCaccaagaggagtagctgcgttcgacagcgacgtgggggagtagcggcggtgacggagctggggcgg  
cctgatgccgagtlactiggaacagccagaaggacctccctggagcagaggcgggcccgggtggacaaTtactgcagac  
acaactacggggttgtGgagagcttcacagtcagc (SEQ ID NO: 90) ;

DRB1\*070101 :

atgggtgctgaagctccctggaggctccctgcatggcagctctgacagtgacactgaatgggtgctgagctccccac  
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gcgggtgcagttccctggaaagactcttctataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagttac  
5 cgggcgggtgacggagctagggcggcctgtcgcgagttccctggaacagccagaaggacatccctggaggacaggcggg  
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NO: 9 1) ;

DRB1\*070102 :

cacgtttccctgtggcagggttaaAtataaagtgctatttcttcaacgggacggagcgggtgcagttccctggaaagact  
10 cttctataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagttaccgggcgggtgacggagctAgggcgg  
cctgtcgcgagttcctggaacagccagaaggacatccctggaggacaggcggggccagggtggacaccgtGtgcagac  
acaactacgggggttggtg (SEQ ID NO: 9 2) ;

DRB1\*0703 :

cacgtttccctgtggcagggttaagtataaagtgctatttcttcaacgggacggagcgggtgcagttccctggaaagTct  
15 cttctataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagttaccgggcgggtgacggagctAgggcgg  
cctgtcgcgagttccctggaacagccagaaggacatccctggaggacaggcggggccagggtggacaccgtgtgcagac  
acaactacgggggttggtg (SEQ ID NO: 9 3) ;

DRB1\*0704 :

tttccctgtggcagggttaagtataaagtgctatttcttcaacgggacggagcgggtgcagttccctggaaagactcttc  
20 tataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagttaccgggcgggtgacggagctAgggcggcctg  
tcgccgagttccctggaacagccagaaggacatccctggaggacaggcggggccagggtggacaaTtactgcagacacaa  
ctacgggggttggtgagagc (SEQ ID NO: 9 4) ;

DRB1\*0705 :

cacgtttccctgtggcagggttaagtataaagtgctatttcttcaacgggacggagcgggtgcagttccctggaaagact  
25 cttctataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagttaccgggcgggtgacggagctAgggcgg  
cctgtcgcgagttccctggaacagccGgaaggacatccctggaggacaggcggggccagggtggacaccgtgtgcagac  
acaactacgggggttggtgagagcttcacag (SEQ ID NO: 9 5) ;

DRB1\*0706 :

cacgtttccgtggcagggttaagtataagtgatcttcttcaacgggacggagcgggtgcagttccitggaaagact  
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cctgtcgcGgagtactggaacagccagaaggacatccitggaggacaggcggggccaggtaggacaccgtGtgcagac  
5 acaactacggggttggtgagagcttcacagtgacggcgag (SEQ ID NO: 96) ;

DRB1\*0707 :

cacgtttccgtggcagggttaagtataagtgatcttcttcaacgggacggagcgggtgcagttccitggaaagact  
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cctgtcgcggagttcctggaacagccagaaggacatccitggaggacaggcggggccaggtaggacaccgtgtgcagac  
10 acaactacggggttggtgagagcttcacagt (SEQ ID NO: 97) ;

DRB1\*080101 :

ggggacacccgaccacgtttcttggagtactctacgggtgagtgTatttcttcaatgggacggagcgggtgcggt  
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ggagctggggcggcctagGccgagtlactggaacagccagaaggacTtcttgaagacaggcggggccTggtggac  
15 acctactgcagacacaactacggggttggtgagagcttcacGgtgcagcgcgag (SEQ ID NO: 98) ;

DRB1\*080102 :

cacgtttcttggagtactctacgggtgagtgTatttcttcaatgggacggagcgggtgcggttccitggacagata  
Ttctataaccaagaggagtlacgtcgcttcgacagcgacgtgggggagtlaccgggcggtagcggagctggggcgg  
cctagcggcgagtlactggaacagccagaaggacttcttgaagacaggcggggccitggtagcaccitactgcagac  
20 acaactacggggttggtgagagcttcacggtgcagcgcgag (SEQ ID NO: 99) ;

DRB1\*080201 :

atggtgtgtctgaggctcccitggaggctcttgcattggcagTctgcagtgacactgatgggtgtgagctcccccac  
tggctttggctggggacaccagaccagtttcttggagtactctacgggtgagtgTatttcttcaatgggacgga  
gcgggtgcggttccitggacagatacttctataaccaagaggagtlacgtcgcttcgacagcgacgtgggggagtlac  
25 cgggcggtagcggagctggggcggccitgctccgagtlactggaacagccagaaggacttcttgaagacaggcggg  
cccitggtagcaccitactgcagacacaactacggggttggtgagagcttcacGgtgcagcgcgag (SEQ ID  
NO: 100) ;

DRB1\*080202 :

cacgtttcttggagtactctacgggtgagtgTattttcttcaatgggacggagcgggtgcggttccitggacagata  
cttctataaccaagaggagTAcgtgcgcttcgacagcgacgtgggggagTaccgggcggtgacggagctggggcgg  
cctgatgccgagTacttgaacagccagaaggacTtccitggaagacaggcggggcccTggtggacacctactgcagac  
5 acaactacggggttggtagagcttcacagtgcagcggcgag (SEQ ID NO: 1 0 1) ;

DRB1\*080203 :

cgtttcttggagtactctacgggtgagtgTattttcttcaatgggacggagcgggtgcggttccitggacagatact  
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tgatgccgagTacttgaacagccagaaggacTtccitggaagacaggcggggcccTggtggacacctactgcagacac  
10 aactacggggttggtagagcttcacgggtg (SEQ ID NO: 1 0 2) ;

DRB1\*080302 :

ggggacaccagaccacgtttcttggagtactctacgggtgagtgTattttcttcaatgggacggagcgggtgcggt  
tccitggacagatactttctataaccaagaggagTactgtgcgcttcgacagcgacgtgggggagTaccgggcggtgac  
ggagctggggcggcctGcgccgagTacttgaacagccagaaggacTtccitggaagacaggcggggcccTggtggac  
15 acctactgcagacacaactacggggttggtagagcttcacagtgcagcggcgag (SEQ ID NO: 1 0 3) ;

DRB1\*080401 :

ggggacaccagaccacgtttcttggagtactctacgggtgagtgTattttcttcaatgggacggagcgggtgcggt  
tccitggacagatactttctataaccaagaggagTAcgtgcgcttcgacagcgacgtgggggagTaccgggcggtgac  
ggagctggggcggcctgatgccgagTacttgaacagccagaaggacTtccitggaagacaggcggggcccTggtggac  
20 acctactgcagacacaactacggggttggtagagcttcacagtgcagcggcgag (SEQ ID NO: 1 0 4) ;

DRB1\*080402 :

ttcaatgggacggagcgggtgcggttccitggacagatactttctataaccaagaggagTAcgtgcgcttcgacagcg  
acgtgggggagTaccgggcggtgacggagctggggcggcctgatgccgagTacttgaacagccagaaggacTtccit  
ggaagacaggcggggcccTggtggacacctactgcagacacaactacggggttggtagagcttcacagtgcagcgg  
25 (SEQ ID NO: 1 0 5) ;

DRB1\*080403 :

cacgtttcttggagtactctacgggtgagtgTattttcttcaatgggacggagcgggtgcggttccitggacagata

cttctataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagtaccgggcggtgacggagctggggcgg  
cctgatgccgagtagtggaaacagccagaaggacttccctggaagacaggcgggcccTggtggacacctactgcagac  
acaactacggggttgTigagagcttcacGgtgcagcggcgag (SEQ ID NO: 106) ;

DRBI\*080404 :

- 5 cactttcttggagtagtctacgggtgagtgtaatttcttcaatgggacggagcgggtgcggttccctggacagata  
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cctgatgccgagtagtggaaacagccagaaggacttccctggaagacaggcgggcccTggtggacacctactgcagac  
acaactacggggttgTigagagcttcacGgtgcagcggcgag (SEQ ID NO: 107) ;

DRBI\*0805 :

- 10 cactttcttggagtagtctacgggtgagtgtaatttcttcaatgggacggagcgggtgcggttccctggacagata  
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cctaGcgccgagtagtggaaacagccagaaggacttccctggaagaCaggcgggcccgcggtggacacctactgcagac  
acaactacggggttggt (SEQ ID NO: 108) ;

DRBI\*0806 :

- 15 ccactttcttggagtagtctacgggtgagtgtaatttcttcaatgggacggagcgggtgcggttccctggacagat  
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gcctaGcgccgagtagtggaaacagccagaaggacttccctggaagacaggcgggcccTggtggacacctactgcaga  
cacaactacggggttgTigagagcttcacagtgacggcgag (SEQ ID NO: 109) ;

DRBI\*0807 :

- 20 cactttcttggagtagtctacgggtgagtgtaatttcttcaatgggacggagcgggtgcggttccctggacagata  
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cctgTlgccgagtagtggaaacagccagaaggacttccctggaagacaggcgggcccTggtggacacctactgcagac  
acaactacggggttggtgagagcttcacGgtgcagcggcgag (SEQ ID NO: 110) ;

DRBI\*0808 :

- 25 tggagtagtctacgggtgagtgtaatttcttcaatgggacggagcgggtgcggttccctggacagatacttctata  
accaagaggagtacgtgcgcttcgacagcgacgtgggggagtaccgggcggtgacggagctggggcggcctgCtgc  
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ggggttggtag(SEQ ID NO: 1 1 1) ;

DRB1\*0809 :

cacgtttcttggagtactctacgggtgagtggtatttcttcaatgggacggagcgggtgcggttcttggacagata  
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5 cctgatgccgagtactggaacagccagaaggacTtccgtggaagacaggcgggcccTggtggacacctactgcagac  
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DRB1\*0810 :

cacgtttcttggagtactctacgggtgagtggtatttcttcaatgggacggagcgggtgcggttcttggacagata  
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10 cctacGcgccgagtactggaacagccagaaggacAtcctggaagacaggcgggcccTggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtgacggcgag(SEQ ID NO: 1 1 3) ;

DRB1\*0811 :

cacgtttcttggagtactctacgggtgagtggtatttcttcaatgggacggagcgggtgcggttcttggacagata  
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15 cctgCtgccgagtactggaacagccagaaggacttccgtggaagacaggcgggcccTggtggacacctactgcagac  
acaactacggggttggtagagcttcacGgtg(SEQ ID NO: 1 1 4) ;

DRB1\*0812 :

cacgtttcttggagtactctacgggtgagtggtatttcttcaatgggacggagcgggtgcggttcttggacagata  
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20 cctagcgccgagtactggaacagccagaaggacAtcctggaagacaggcgggcccTggtggacacctactgcagac  
acaactacggggCgtggagagcttcacagtgacggcgag(SEQ ID NO: 1 1 5) ;

DRB1\*0813 :

tcttggagtactctacgggtgagtggtatttcttcaatgggacggagcgggtgcggttcttggacagatacttcta  
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25 gccgagtactggaacagccagaaggaccttccgtggaagacaggcgggcccTggtggacacctactgcagacacaact  
acggggttggtagagcttcacGgtg(SEQ ID NO: 1 1 6) ;

DRB1\*0814 :

cacgtttcttggagtactcttaGgggtgagtggtattttcttcaatgggacggagcgggtgcggttccctggacagata  
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cctagcgccgagtagctggaacagccagaaggacatcttgaagacaggcggggccctggtagcacctactgcagac  
acaactacggggttggtagagcttcacagtg (SEQ ID NO: 117) ;

5 DRB1\*0815 :

tttcttggagtactctacgggtgagtggtattttcttcaatgggacggagcgggtgcggttccctggacagatacttc  
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atgcggagCactggaacagccagaaggacAtcctggaagacaggcggggccctTggtagcacctactgcagacacaa  
ctacggggttggtag (SEQ ID NO: 118) ;

10 DRB1\*0816 :

cacgtttcttggagtactctacgggtgagtggtattttcttcaatgggacggagcgggtgcggttccctggacagata  
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cctagcgccgagtagctggaacagccagaaggacttcttgaagacaggcggggccctggtagcacctactgcagac  
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15 DRB1\*0817 :

cacgtttcttggagtactctacgggtgagtggtattttcttcaatgggacggagcgggtgcggttccctggacagata  
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cctaGcgccgagtagctggaacagccagaaggacTtcttgaagacaggcggggccctTggtagcacctactgcagac  
acaactacggggttggtag (SEQ ID NO: 120) ;

20 DRB1\*0818 :

cacgtttcttggagtactcttaCgggtgagtggtattttcttcaatgggacggagcgggtgcggttccctggacagata  
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cctaGcgccgagtagctggaacagccagaaggacAtcctggaagaCaggcggggcccggttagcacctactgcagac  
acaactacggggttggtagagcttcacagtagcagcggcgag (SEQ ID NO: 121) ;

25 DRB1\*0819 :

tttcttggagtactctacgggtgagtggtattttcttcaatgggacggagcgggtgcggttccctggacagatacttc  
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TcgccgagtlactggaacagccagaaggacAtcctggaagacaggcgggccTggtggacacctactgcagacacaa  
ctacggggttggtgagagcttcacagtgc (SEQ ID NO: 1 2 2) ;

DRB1\*0820 :

cacgtttcttggagtlactctacgtCtgagtgtcatttcttcaatgggacggagcgggtgcggttccctggacagata  
5 cttctataaccaagaggagtAcgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
ccgatgccgagtlactggaacagccagaaggacTtcttgaagacaggcgggccTggtggacacctactgcagac  
acaactacggggttGgagagcttcacagtcagcggcga (SEQ ID NO: 1 2 3) ;

DRB1\*0821 :

cacgtttcttggagtlactctatgggtgagtgtaatttcttcaatgggacggagcgggtgcggttccctggacagata  
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cctgatgccgagtlactggaacagccagaaggacttcttgaagacaggcgggccctgggtggacacctactgcagac  
acaactacggggttggtagagcttcacggtgcagcggcga (SEQ ID NO: 1 2 4) ;

DRB1\*0822 :

cacgtttcttggagtlactctacgggtgagtgtaatttcttcaatgggacggagcgggtgcggttccctggacagata  
15 cttctataaccaagaggagtlacgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
cctagcggcagtlactggaacagccagaaggacttcttgaagacaggcgggccctgggtggacacctactgcagac  
acaactacggggCgtGgagagcttcacGgtgcagcggcgag (SEQ ID NO: 1 2 5) ;

DRB1\*0823 :

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20 cttctataaccaagaggagtlacgtgcgcttcgacagcgacgtAgggagtlaccgggcggtgacggagctggggcgg  
cctagcggcagtlactggaacagccagaaggacatcttgaagacaggcgggccTggtggacacctactgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 1 2 6) ;

DRB1\*0824 :

cacgtttcttggagtlactctacgggtgagtgtaatttcttcaatgggacggagcgggtgcggttccctggacagata  
25 cttctataaccaagaggagtAcgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
ccgatgccgagtlactggaacagccagaaggacTtcttgaagaCaggcgggcccggtggacacctactgcagac  
acaactacggggttggtagagcttcacagtcagcgg (SEQ ID NO: 1 2 7) ;

DRB1\*090102 :

ggggacacccaaccacgtttcttgaagcaggataagtttgagtgtcatitcttcaacgggacggagcgggtgcggt  
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5 accgtgtgcagacacaactacggggttggtgagagcttcacagtgcagAggcgag (SEQ ID NO: 1 2 8) ;

DRB1\*0902 :

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DRB1\*100101 :

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15 cgggcggtgacggagctggggcggccctgatcccgagtagcgaacagccagaaggacctccctggagcggaggcgTg  
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NO: 1 3 0) ;

DRB1\*100102 :

cacgtttcttggaggaggtaagtttgagtgtcatitcttcaacgggacggagcgggtgcggttgcctggaagacg  
20 cGtccataaccaagaggagtagcgcgctacgacagcgacgtgggggagtagcgggcggtgacggagctggggcgg  
ccctgatcccgagtagcgaacagccagaaggacctccctggagcggaggcgGccgcggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtgcagcggcgag (SEQ ID NO: 1 3 1) ;

DRB1\*110101 :

atgggtgtgtctgaggctccctggaggctccctgcatggcagTtctgacagtgcacacgatgggtgtgagctccccac  
25 tggccttggctggggacaccagaccacgtttcttggagtactctacgtctgagtgtcatitcttcaatgggacgga  
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ccgcgggtggacacctactgcagacacaactacgggggtggtagagcttcacagtcagcggcgag (SEQ ID NO: 1 3 2) ;

DRB1\*110102 :

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5 tcttggacagatacttctataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagttccgggcggtgac  
ggagctggggcggccctgatgaGgagtactggaacagccagaaggacTtccctggaagaCaggcgggcccgcggtggac  
acctactgcagacacaactacgggggtggtagagcttcacGgtgcagcggcgag (SEQ ID NO: 1 3 3) ;

DRB1\*110103 :

cacgtttcttggagtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttcttggacagata  
10 ctctataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
ccctgatgaGgagtactggaacagccagaaggacTtccctggaagaCaggcgCgccgcggtggacacctactgcagac  
acaactacgggggtggtagagcttcacagtcagcggcgag (SEQ ID NO: 1 3 4) ;

DRB1\*110104 :

cgtttcttggagtactctacgtctgagtgctcatttcttcaaCgggacggagcgggtgcggttcttggacagatact  
15 tctataaccaagaggagtAcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcc  
tgatgAggagtactggaacagccagaaggacTtccctggaagaCaggcgggcccgcggtggacacctactgcagacac  
aactacgggggtggtagagcttcacagtcagcggcgag (SEQ ID NO: 1 3 5) ;

DRB1\*1102 :

ggggacaccagaccacgtttcttggagtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggt  
20 tcttggacagatacttctataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagttccgggcggtgac  
ggagctggggcggccctgatgAggagtactggaacagccagaaggacAtccctggaagacGAgcgggcccgcggtggac  
acctactgcagacacaactacgggggtgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 1 3 6) ;

DRB1\*1103 :

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25 tggctttggctggggacaccagaccacgtttcttggagtactctacgtctgagtgctcatttcttcaatgggacgga  
gcgggtgcggttcttggacagatacttctataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagttc  
cgggcggtgacggagctggggcggccctgatgaggagtactggaacagccagaaggacTtccctggaagacGAgcggg

ccgcggiggacacctactgcagacacaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID  
NO: 1 3 7) ;

DRB1\*110401 :

atgggtgigtctgaggctccctggaggctcctgcatggcagTcttgacagtgacactgatgggtgctgagctccccac  
5 tggctttggctggggacaccagaccacgtttcttggagtactctacgtctgagtgctatitcttcaatgggacgga  
gcgggtgaggcttctggacagatctctataaccaagaggagtagctgcgcttcgacagcgacgtgggggagttc  
cgggcggtagcggagctggggcggcctgatgaggagtagtgaacagccagaaggacTtcttgaagaCaggcggg  
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NO: 1 3 8) ;

10 DRB1\*110402 :

ggggacaccagaccacgtttcttggagtactctacgtctgagtgctatitcttcaatgggacggagcgggtgagg  
tcttggacagatacttctataaccaagaggagtagctgcgcttcgacagcgacgtgggggagttccgggcggtag  
ggagctggggcggcctgatgaGgagtactggaacagccagaaggacttcttgaagacaggcgggcccgggtggac  
acctactgcagacacaactacggggttgtGgagagcttcacGgtgcagcggcgag (SEQ ID NO: 1 3 9) ;

15 DRB1\*1105 :

ccacgtttcttggagtactctacgtctgagtgctatitcttcaatgggacggagcgggtgaggcttcttggacagat  
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gcctgatgAGgagtactggaacagccagaaggacTtcttgaagaCaggcgggcccgggtggacacctactgcaga  
cacaactacggggttggtagagcttcacagtcagcggcgga (SEQ ID NO: 1 4 0) ;

20 DRB1\*110601 :

cgtttcttggagtactctacgtctgagtgctatitcttcaatgggacggagcgggtgaggcttcttggacagatact  
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tgatgaGgagtactggaacagccagaaggacTtcttgaagaCaggcgggcccgggtggacacctactgcagacac  
aactacggggCtggtagagcttcacagtcagcggcgag (SEQ ID NO: 1 4 1) ;

25 DRB1\*110602 :

tttcttggagtactctacgtctgagtgctatitcttcaatgggacggagcgggtgaggcttcttggacagatactt  
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atgaGgagtaciggaacagccagaaggacTtccctggaagaCaggcgggccggttgacacctatTgcagacacaa  
ctacggggCtctggagagcttcacagtcagcggcgag (SEQ ID NO: 1 4 2) ;

DRB1\*1107 :

ttggagtactctacgtctgagtgctatttcttcaatgggacggagcgggtgcggttccctggacagatacttctata  
5 accaagaggagtacgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgatgA  
GgagtactiggaacagccagaaggaccttctggagcagaagcggggccGggtggacaActactgcagacacaactac  
ggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 1 4 3) ;

DRB1\*110801 :

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10 gtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgatgAggagtactiggaaca  
gccagaaggaccttctggaagaCaggcgggccggttgacacctactgcagacacaactacggggttggtgagag  
cttcacagtg (SEQ ID NO: 1 4 4) ;

DRB1\*110802 :

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15 gtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgatgGgagtactiggaaca  
gccagaaggaccttctggaagaCaggcgggccggttgacacctactgcagacacaactacggggttggtgagag  
cttcacGgtg (SEQ ID NO: 1 4 5) ;

DRB1\*1109 :

catttcttcaatgggacggagcgggtgcggttccctggacagatacttccataaccaGaggagAAcgtgcgcttcg  
20 acagcgacgtgggggagttccgggcggtgacggagctggggcggcctgatgAggagtactiggaacagccagaagga  
cTtccctggaagaCaggcgggccggttgacacctactgcagacacaactacggggttggtgagagcttcacagtg  
cag (SEQ ID NO: 1 4 6) ;

DRB1\*1110 :

gagtgctatttcttcaatgggacggagcgggtgcggttccctggacagatacttccataaccaGaggagtTcgtgc  
25 gcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgatgAggagtactiggaacagcca  
gaaggacTtccctggaagaCaggcgggccggttgacacctactgcagacacaactacggggttggt (SEQ ID  
NO: 1 4 7) ;

DRB1\*1111 :

tttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggttccctggacagatacttc  
tataaccaagaggagtacgtgcgttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcgccctg  
atgAggagtiactggaacagccagaaggacTtccctggaagacGAgcgggccggtggacacctactgcagacacaa  
5 ctacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 148) ;

DRB1\*111201 :

gagtgatcatttcttcaatgggacggagcgggtgcggttccctggacagatacttctataaccaagaggagtTcgtgc  
gcttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcgccctgatgAggagtiactggaacagcca  
gaaggacTtccctggaagaCaggcgggccggtggacacctactgcagacacaactacggggttggt (SEQ ID  
10 NO: 149) ;

DRB1\*111202 :

cacgtttcttggagtactctacgtCtgagtgatcatttcttcaatgggacggagcgggtgcggttccctggacagata  
cttctataaccaGgaggagtTcgtgcgttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcgcc  
cctgatgAggagtiactggaacagccagaaggacTtccctggaagaCaggcgggccggtggacacctactgcagac  
15 acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 150) ;

DRB1\*1113 :

ggggacaccagaccacgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggt  
tccctggacagatacttccataaccaggaggagttcgtgcgttcgacagcgacgtgggggagtTccgggcggtgac  
ggagctggggcgccctgatgAGgagtiactggaacagccagaaggaccttctggagcGgaggcgggccggtggac  
20 acciaTtgcagacacaactacggggttgtGgagagcttcacagtcagcggcgga (SEQ ID NO: 151) ;

DRB1\*1114 :

ggggacaccagaccacgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggt  
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ggagctggggcgccctgatgAggagtiactggaacagccagaaggacAtcctggaagacGAgcgggccggtggac  
25 acctactgcagacacaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 152) ;

DRB1\*1115 :

ggggacaccagaccacgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggt



tccitggacagatacttctataaccaagaggaggacTtgcgcttcgacagcgacgtgggggagttccgggcgggtgac  
ggagctggggcggcctgatgaGgagtacttggaacagccagaaggacTtccitggaaGaCaggcgggcccggtggac  
acctactgcagacacaacttacggggttggtgagagcttcacagtgcagcggcgag (SEQ ID NO: 1 5 3) ;  
DRB1\*1116 :

5 cactttctitggagtactctacgtctgagtgtcatttcttcaatgggacggagcgggtgcggttccitggacagata  
cttccataaaccaggaggagAacgtgcgcttcgacagcgacgtgggggagttccgggcgggtgacggagctggggcgg  
cctgatgAggagtacttggaacagccagaaggacAtccttggaagacGAgcgggcccggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 1 5 4) ;  
DRB1\*1117 :

10 ggggacaccagaccacgtttctitggagtactctacgtCtgagtgtcatttcttcaatgggacggagcgggtgcggt  
tccitggacagatacttccataaaccaggaggagttcgtgcgcttcgacagcgacgtgggggagttaccgggcgggtgac  
ggagctggggcggcctgatgAGgagtacttggaacagccagaaggacctccitggagcggaggcgggccgAggtggac  
acctatTgcagacacaacttacggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 1 5 5) ;  
DRB1\*1118 :

15 tttctitggagtactctacgtctgagtgtcatttcttcaatgggacggagcgggtgcggttccitggacagatacttc  
tataaccaagaggagtAcgtgcgcttcgacagcgacgtgggggagttccgggcgggtgacggagctggggcggcctg  
atgAggagtacttggaacagccagaaggacAtccttggaagaCaggcgggcccggtggacacctactgcagacacaa  
ctacggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 1 5 6) ;  
DRB1\*1119 :

20 tttctitggagtactctacgtctgagtgtcatttcttcaatgggacggagcgggtgcggttccitggacagatacttc  
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atgAggagtacttggaacagccagaaggacAtccttggaagaCaggcgggcccggtggacacctactgcagacacaa  
ctacggggttggtgagagcttcacagtgcagcggcgag (SEQ ID NO: 1 5 7) ;  
DRB1\*1120 :

25 ttggagtactctacgtctgagtgtcatttcttcaatgggacggagcgggtgcggttccitggacagatacttccata  
accaggaggagAacgtgcgcttcgacagcgacgtgggggagttccgggcgggtgacggagctggggcggcctgatgA  
ggagtacttggaacagccagaaggacAtccttggaagacGAgcgggcccggtggacacctactgcagacacaacttac

ggggttggtgagagcttcacagtcagc (SEQ ID NO: 1 5 8) ;

DRB1\*1121 :

ttggagtlactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagatcttctata  
accaagaggagtlactgtcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgccgatga  
5 ggagtlactggaacagccagaaggacatcctggaagacGAgcgggcccgggtggacacctactgcagacacaactac  
ggggCtgggaga (SEQ ID NO: 1 5 9) ;

DRB1\*1122 :

cacgtttcttggagcaggtttaaCatgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagata  
cttctataaaccaggaggagtlactgtcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcg  
10 cctgatgaGgagtlactggaacagccagaaggacTtcttgaagaCaggcgggcccgggtggacacctactgcagac  
acaactacggggttggtagag (SEQ ID NO: 1 6 0) ;

DRB1\*1123 :

ccacgtttcttggagtlactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagat  
acttctataaaccaagaggagtlactgtcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcg  
15 gcctgatGAggagtlactggaacagccagaaggacTtcttgaagaCaggcgggcccTggtggacacctactgcaga  
cacaactacggggttggtag (SEQ ID NO: 1 6 1) ;

DRB1\*1124 :

ttcttggagtlactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagatcttct  
ataaccaagaggagGacgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgcciga  
20 tgaGgagtlactggaacagccagaaggacTtcttgaGaCaggcgggcccgggtggacacctactgcagacacaac  
tacggggttggtagagcttcac (SEQ ID NO: 1 6 2) ;

DRB1\*1125 :

cacgtttcttggagtlactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagata  
cttctataaaccaagaggagtlactgtcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcg  
25 cctgatgaGgagtlactggaacagccagaaggacTtcttgaagacaggcgggcccTggtggacacctactgcagac  
acaactacggggttGgagagcttcacagtcagcggcgag (SEQ ID NO: 1 6 3) ;

DRB1\*1126 :

tiggagtlactctacgtCtgaGigtcatttcttcaatgggacggagcgggtgcggttccitggacagatacttctata  
accaagaggagtAcgtgcgcttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcggcctgatGA  
ggagtlacttggaacagccagaaggacctccitggagcagaggcgggccggtggacacctactgcagacacaacttac  
ggggttgggtgag (SEQ ID NO: 164) ;

5 DRB1\*112701 :

tttcttggagtlactctacgtctgagtgctatttcttcaatgggacggagcgggtgcggttccitggacagatacttc  
tataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctg  
atgaggagtlacttggaacagccagaaggacctccitggaAgaCaggcgggccggtggacaaTtactgcagacacaa  
ctacggggttgggtgag (SEQ ID NO: 165) ;

10 DRB1\*112702 :

cacgtttcttggagtlactctacgtctgagtgctatttcttcaatgggacggagcgggtgcggttccitggacagata  
cttctataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
cctgatGAggagtlacttggaacagccagaaggacctccitggaagaCaggcgggccggtggacaaTtactgcagac  
acaactacggggttgggtgagagcttcacagtcagcggcgag (SEQ ID NO: 166) ;

15 DRB1\*1128 :

cacgtttcttggagtlactctacgtctgaGigtcatttcttcaatgggacggagcgggtgcggttccitggacagata  
cttctataaccaagaggagAAcgtgcgcttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcgg  
cctgatGAggagtlacttggaacagccagaaggacTtccitggaagaCaggcgggccggtggacacctactgcagac  
acaactacggggttgggtgagagcttcacagtcagcggcgag (SEQ ID NO: 167) ;

20 DRB1\*1129 :

cacgtttcttggagtlactctacgtctgaGigtcatttcttcaatgggacggagcgggtgcggttccitggacagata  
cttctataaccaagaggagtcggtgcgcttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcgg  
cctgatGAggagtlacttggaacagccagaaggacTtccitggaagaCaggcgggccggtggacacctactgcagac  
acaactacggggttgggtgagagcttcacagtcagcggcgag (SEQ ID NO: 168) ;

25 DRB1\*1130 :

cacgtttcttggagcTgcttaagtcgagtgctatttcttcaatgggacggagcgggtgcggttccitggacagata  
cttctataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg

cctgatgaggagctacaggacagccagaaggacttcctggaagaCaggcgggccggtggacacctactgcagac  
acaactacggggttggtagagcttcacagtcagcggcga (SEQ ID NO: 169) ;

DRB1\*1131 :

ggggacaccagaccagtttcttggagtacctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggt  
5 tcttggacagatacttctataaccaagaggagctacgtgcgcttcgacagcgacgtgggggagttccgggcggtagc  
ggagctggggcgccctgatgAggagCacttggacagccagaaggacAtcctggaagaCaggcgggccggtggac  
acctactgcagacacaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 170) ;

DRB1\*1132 :

cacgtttcttggagtacctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttcttggacagata  
10 ctctataaccaagaggagctacgtgcgcttcgacagcgacgtgggggagttccgggcggtagcggagctggggcgg  
cctgatgAGgagtacttggacagccagaaggacttcctggaagaCaggcgggccgTggtggacacctactgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 171) ;

DRB1\*1133 :

ttggagtacctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttcttggacagatacttctata  
15 accaagaggagctacgtgcgcttcgacagcgacgtgggggagttccgggcggtagcggagctggggcgccctgatga  
ggaCtacttggacagccagaaggacttcctggaagacaggcgggccggtggacacctactgcagacacaactac  
ggggttggtagagcttcacagtcagcggc (SEQ ID NO: 172) ;

DRB1\*1134 :

cacgtttcttggagtacctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttcttggacagata  
20 ctctataaccaagaggagTcgtgcgcttcgacagcgacgtgggggagTccgggcggtagcggagctggggcgg  
cctgatgAggagtacttggacagccagaaggaccttcctggagcagaggcgggccggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcga (SEQ ID NO: 173) ;

DRB1\*1135 :

ttggagtacctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttcttggacagatacttctata  
25 accaagaggagctacgtgcgcttcgacagcgacgtgggggagttccgggcggtagcggagctggggcgccctgatga  
ggaCtacttggacagccagaaggacttcctggaagacaggcgggccggtggacacctactgcagacacaactac  
ggggttgtGgagagcttcacagtcagcggc (SEQ ID NO: 174) ;

DRB1\*1136 :

cgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagatact  
cttataaccaagaggagtagctgcgtctcgacagcgacgtgggggagTccgggcggtgacggagctggggcggcc  
tgatgAggagtagtggaacagccagaaggacctcttggaagacGAgcgggccgcggtggacacctactgcagacac  
5 aactacggggttgtGgagagcttcacagtcagcggcga (SEQ ID NO: 1 7 5) ;

DRB1\*1137 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagata  
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ccctgatgAggagtagtggaacagccagaaggacTtcttggaagaCaggcgggccgcggtggacacctactgcagac  
10 acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 1 7 6) ;

DRB1\*1138 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagata  
cttctataaccaagaggagtagctgcgtctcgacagcgacgtgggggagTccgggcggtgacggagctggggcgg  
ccctgatgaggGgtacttggaacagccagaaggaccttcttggaagacaggcgggccgcggtggacacctactgcagac  
15 acaactacggggttgtggagagcttcacagtcagcggcgag (SEQ ID NO: 1 7 7) ;

DRB1\*1139 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagata  
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ccctgatgaggagtagtggaacagccagaaggaccttcttggaagaCaggcgggccgcggtggacacctactgcagac  
20 acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 1 7 8) ;

DRB1\*1140 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagata  
cttccataaccaggaggagAacgtgcgtctcgacagcgacgtgggggagTccgggcggtgacggagctggggcgg  
ccctgatgAggagtagtggaacagccagaaggacTtcttggaagacGAgcgggccgcggtggacacctactgcagac  
25 acaactacggggttgtGg (SEQ ID NO: 1 7 9) ;

DRB1\*1141 :

tttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagatacttc

tataaccaagaggagtagctgcgcttcgacagcgacgtgggggagtagcgggcggtgacggagctggggcgccctg  
atgAggagtagctggaacagccagaaggacTtccctggaagacGAgcgggcccgcggtggacacctactgcagacacaa  
ctacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 180) ;

DRB1\*1142 :

- 5 cacttttcttgagtagctctacgtctgagtgicatttcttcaatgggacggagcgggtgcggttccctggacagata  
cttctataaccaagaggagtagctgcgcttcgacagcgacgtgggggagtagcgggcggtgacggagctggggcgg  
ccgatgAggagtagctggaacagccagaaggaccttctggaagaCaggcgggcccgcggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 181) ;

DRB1\*1143 :

- 10 cacttttcttgagtagctctacgtctgagtgicatttcttcaatgggacggagcgggtgcggttccctggacagata  
cttctataaccaagaggagtagctgcgcttcgacagcgacgtgggggagtagcgggcggtgaGggagctggggcgg  
ccgatgaggagtagctggaacagccagaaggaccttctggaagaCaggcgggcccgcggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 182) ;

DRB1\*120101 :

- 15 atgggtgtgtctgaggctccctggaggctcctgcatggcagTctgacagtgacactgatgggtgtgagctccccac  
tggctttggctggggacaccAgaccacgtttcttgagtagctctacgggtgagtggtatttcttcaatgggacgga  
gcgggtgcggttActggagagacacttccataaccaggaggagCtccctgcgcttcgacagcgacgtgggggagttc  
cgggcggtgacggagctggggcgccctgtcgccgagTctggaacagccagaaggacAtcctggaagacaggcgcg  
ccgcggtggacacctatTgcagacacaactacgggCtgggagagcttcacagtcagcggcgag (SEQ ID

- 20 NO: 183) ;

DRB1\*120102 :

- atgggtgtgtctgaggctccctggaggctcctgcatggcagTctgacagtgacactgatgggtgtgagctccccac  
tggctttggctggggacaccAgaccacgtttcttgagtagctctacgggtgagtggtatttcttcaatgggacgga  
gcgggtgcggttActggagagacacttccataaccaggaggagCtccctgcgcttcgacagcgacgtgggggagttc  
25 cgggcggtgacggagctggggcgccctgtcgccgagTctggaacagccagaaggacAtcctggaagacaggcggg  
ccgcggtggacacctactgcagacacaactacgggCtgggagagcttcacagtcagcggcgag (SEQ ID

NO: 184) ;

DRB1\*120201 :

cacgtttcttggagtaactctacgggtgagtggtattttcttcaatgggacggagcgggtgcggttactggagagaca  
cttccataaccaggaggagCtccctgcgttcgacagcgacgtgggggagttccgggcgggtgacggagctggggcgg  
ccctgcgccgagtcctggaacagccagaaggacTtccctggaagacaggcgcccgcggtggacacctatTgcagac  
5 acaactacggggCtgtggagagcttcacagtgacggcgag (SEQ ID NO: 185) ;

DRB1\*120202 :

ttcttggagtaactctacgggtgagtggtattttcttcaatgggacggagcgggtgcggttactggagagacacttcc  
ataaccaggaggagCtccctgcgttcgacagcgacgtgggggagttccgggcgggtgacggagctggggcggcctgt  
cgccgagtcctggaacagccagaaggacTtccctggaagacaggcgCgcccgggtggacacctactgcagacacaac  
10 tacggggCtgtggag (SEQ ID NO: 186) ;

DRB1\*120302 :

cacgtttcttggagtaactctacgggtgagtggtattttcttcaatgggacggagcgggtgcggttActggagagaca  
cttccataaccaggaggagCtccctgcgttcgacagcgacgtgggggagttccgggcgggtgacggagctggggcgg  
ccctgcgccgagtcctggaacagccagaaggacAtcctggaagacaggcgCgcccgggtggacacctactgcagac  
15 acaactacggggltgtggagagcttcacagtgacgg (SEQ ID NO: 187) ;

DRB1\*1204 :

gagtaactctacgggtgagtggtattttcttcaatgggacggagcgggtgcggttactggagagacacttccataacc  
aggaggagCtccctgcgttcgacagcgacgtgggggagttccgggcgggtgacggagctggggcggcctgataGga  
gtactggaacagccagaaggacAtcctggaagacaggcgcccgcggtggacacctatTgcagacacaactacggg  
20 gCtgtgg (SEQ ID NO: 188) ;

DRB1\*1205 :

cacgtttcttggagtaactctacgggtgagtggtattttcttcaatgggacggagcgggtgcggttActggagagaca  
cttccataaccaggaggagttcctgcgttcgacagcgacgtgggggagttccgggcgggtgacggagctggggcgg  
ccctgcgccgagtcctggaacagccagaaggacAtcctggaagacaggcgcccgcggtggacacctatTgcagac  
25 acaactacggggCtgtggagagcttcacagtgacggcgag (SEQ ID NO: 189) ;

DRB1\*1206 :

ggggacaccagaccacgtttcttggagtaactctacgggtgagtggtattttcttcaatgggacggagcgggtgcggt

tActggagagacacttccataaccaggaggagCtctgcgcttcgacagcgacgtgggggagttccgggcggtagc  
ggagctggggcggcctgtgccgagtcctggaacagccagaaggacAtcctggaagacaggcgcccgcggtggac  
acctatTgcagacacaacttacggggCtgtggagagcttcacagtcagcggcgag (SEQ ID NO: 190) ;  
DRB1\*1207 :

5 cactttcttggagtactctacgggtgagtggtatttcttcaatgggacggagcgggtgcggttactggagagaca  
cttccataaccaggaggagcttctgcgcttcgacagcgacgtgggggagttccgggcggtagcggagctggggcgg  
cctgtgccgagtcctggaacagccagaaggacatcctggGagacaggcgcccgcggtggacacctatTgcagac  
acaactacggggCtgtggagagcttcacagtcagcggcgag (SEQ ID NO: 191) ;  
DRB1\*1208 :

10 cactttcttggagtactctacgggtgagtggtatttcttcaatgggacggagcgggtgcggttCctggagagaca  
cttccataaccaggaggagCtctgcgcttcgacagcgacgtgggggagttccgggcggtagcggagctggggcgg  
cctgtgccgagtcctggaacagccagaaggacAtcctggaagacaggcgcccgcggtggacacctatTgcagac  
acaactacggggCtgtggagagcttcacagtcagcggcgag (SEQ ID NO: 192) ;  
DRB1\*130101 :

15 ggggacaccagaccacgtttcttggagtactctacgtctgagtgctatttcttcaatgggacggagcgggtgcggt  
tcttgaCagatacttccataaccaggaggagAAcgtgcgcttcgacagcgacgtgggggagttccgggcggtagc  
ggagctggggcggcctgatgccgagtagtgaacagccagaaggacAtcctggaagacAGcggggcccggtggac  
acctactgcagacacaactacggggtgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 193) ;  
DRB1\*130102 :

20 cactttcttggagtactctacgtctgagtgctatttcttcaatgggacggagcgggtgcggttcttgacagata  
cttccataaccaggaggagaacgtgcgcttcgacagcgacgtgggggagttccgggcggtagcggagctggggcgg  
cctgatgccgagtagtgaacagccagaaggacatcctggaagacgagcgggcTgcggtggacacctactgcagac  
acaactacggggtgtggagagcttcacagtcagcggcgag (SEQ ID NO: 194) ;  
DRB1\*130103 :

25 cactttcttggagtactctacgtctgagtgctatttcttcaatgggacggagcgggtgcggttcttgacagata  
cttccataaccaggaggagaacgtgcgcttcgacagcgacgtgggggagttccgggcggtagcggagctggggcgg  
cctgatgccgagtagtgaacagccagaaggacatcctggaagacgAgcggggcccggtggacacctatTgcagac



acaactacggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 195) ;

DRB1\*130201 :

ggggacaccagaccacgtttcttggagtactctacgtctgagtgicatttcttcaatgggacggagcgggtgcggt  
tcttggacagatacttccataaccaggaggagAACgtgcgttcgacagcgacgtgggggagtTccgggcgggtgac  
5 ggagctggggcggcctgatgccgagtlactggaacagccagaaggacAtcctggaagacGAgcgggcccgggtggac  
acctactgcagacacaactacggggttggtagagcttcacagtgcagcggcgag (SEQ ID NO: 196) ;

DRB1\*130202 :

cacgtttcttggagtactctacgtctgagtgicatttcttcaatgggacggagcgggtgcggttcttggacagata  
cttccataaccaggaggagaacgtgcgttcgacagcgacgtgggggagttccgggcgggtgacggagctggggcgg  
10 cctgatgccgagtlactggaacagccagaaggacatcctggaagacgAgcgCgcccgggtggacacctactgcagac  
acaactacggggttggtagagcttcacagtgcagcgg (SEQ ID NO: 197) ;

DRB1\*130301 :

ggggacaccagaccacgtttcttggagtactctacgtctgagtgicatttcttcaatgggacggagcgggtgcggt  
tcttggacagatacttctataaccaagaggagtlactgtgcgttcgacagcgacgtgggggagttaccgggcgggtgac  
15 ggagctggggcggcctagCgccgagtlactggaacagccagaaggacatcctggaagaCaAgcgggcccgggtggac  
acctactgcagacacaactacggggttggtagagcttcacGgtgcagcggcgag (SEQ ID NO: 198) ;

DRB1\*130302 :

tiggagtactctacgtctgagtgicatttcttcaatgggacggagcgggtgcggttcttggacagatacttctata  
accaagaggagtlactgtgcgttcgacagcgacgtgggggagttaccgggcgggtgacggagctggggcggcctaGcgc  
20 cgagtlactggaacagccagaaggacatcctggaagaCaAgcgggcccgggtggacacctactgcagacacaactac  
ggggttggtagagcttcacagtgcagcggcgag (SEQ ID NO: 199) ;

DRB1\*1304 :

ggggacaccagaccacgtttcttggagtactctacgtctgagtgicatttcttcaatgggacggagcgggtgcggt  
tcttggacagatacttctataaccaagaggagtlactgtgcgttcgacagcgacgtgggggagtTccgggcgggtgac  
25 ggagctggggcggcctaGcggcgagtlactggaacagccagaaggacatcctggaagacGAgcgggcccgggtggac  
acctactgcagacacaactacggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 200) ;

DRB1\*1305 :

cgtttcttggagtactctacgtctgaGtgctatttcttcaatgggacggagcgggtgcggttccctggacagatact  
tccataaccaGgaggagAacgtgcgcttcgacagcgacgtgggggagtTccgggcgggtgacggagctggggcggcc  
tgatgccgagtagtggacagccagaaggacTtccctggaagaCaggcgggccgcggtggacacctactgcagacac  
aactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 2 0 1) ;

5 DRB1\*1306 :

tgtcatttcttcaatgggacggagcgggtgcggttccctggaCagatacttccataaccaggaggagAacgtgcgct  
tcgacagcgacgtgggggagtTccgggcgggtgacggagctggggcggcctgatgccgagtagtggacagccagaa  
ggacAtccctggaagaCaggcgggccgcggtggacacctactgcagacacaactacggggttgtGgagagcttcaca  
(SEQ ID NO: 2 0 2) ;

10 DRB1\*130701 :

cacgttcttggagtagtCtaCgtCtgaGtgctatttcttcaatgggacggagcgggtgcggttccctggaCagata  
cTtctataaccaagaggagtAcgtgcgcttcgacagcgacgtgggggagtaccgggcgggtgacggagctggggcgg  
ccctgatgccgagtagtggacagccagaaggacTtccctggaGaCaggcgggccgcggtggacacctactgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 2 0 3) ;

15 DRB1\*130702 :

cacgttcttggagtagtctacgtCtagtgctatttcttcaatgggacggagcgggtgcggttccctggaCagata  
cTtctataaccaagaggagtAcgtgcgcttcgacagcgacgtgggggagtaccgggcgggtgacggagctggggcgg  
ccctgacgtGagtagtagtggacagccagaaggacTtccctggaGaCaggcgggccgcggtggacacctactgcagac  
acaactacggggttggtagagcttcacagtcagcggcg (SEQ ID NO: 2 0 4) ;

20 DRB1\*1308 :

tcttggagtagtctacgtctgagtgctatttcttcaatgggacggagcgggtgcggttccctggaCagatacttcc  
ataaccaggaggagtTcgtgcgcttcgacagcgacgtgggggagtaccgggcgggtgacggagctggggcggcctga  
tgccgagtagtggacagccagaaggacatccctggaagacGAgcgggccgcggtggacacctactgcagacacaac  
tacggggttgtGgagagcttcacagt (SEQ ID NO: 2 0 5) ;

25 DRB1\*1309 :

tcttggagtagtctacgtctgagtgctatttcttcaatgggacggagcgggtgcggttccctggacagatacttc  
cataaccaggaggagaAcgtgcgcttcgacagcgacgtgggggagtTccgggcgggtgacggagctggggcggcctg

atgccgagtlactggaacagccagaaggacAtcctggagcaggCgcgggccgcggtggacacctactgcagacacaa  
ctacgggggttgtGgagagcttcacagt(SAQ ID NO: 2 0 6) ;

DRB1\*1310 :

cacgtttcttggagtlactctacgtctgagtgcatcttctcaatgggacggagcgggtgcggttccctggaCagata  
5 cttccataaaccaggaggagAACgtgcgttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcgg  
ccgatgccgagtlactggaacagccagaaggacAtcctggaagaCaAgcgggccgcggtggacacctactgcagac  
acaactacgggggttgtGgagagcttcacagtcagcggcgag(SAQ ID NO: 2 0 7) ;

DRB1\*1311 :

cacgtttcttggagtlactctacgtctgagtgcatcttctcaatgggacggagcgggtgcggttccctggacagata  
10 cttctataaccaagaggagtAcgtgcgttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcgg  
ccgatgccgagtlactggaacagccagaaggacTtcttggagaCaggcgggccgcggtggacacctactgcagac  
acaactacgggggttgtGgagagcttcacagtcagcggcgag(SAQ ID NO: 2 0 8) ;

DRB1\*1312 :

cacgtttcttggagtlactctacgtctgagtgcatcttctcaatgggacggagcgggtgcggttccctggacagata  
15 cttctataaccaagaggagtacgtgcgttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
cctaGcgccgagtlactggaacagccagaaggacAtcctggaagaCaggcgggccgcggtggacacctactgcagac  
acaactacgggggttggtgagagcttcacagtcagcggcgag(SAQ ID NO: 2 0 9) ;

DRB1\*1313 :

cacgtttcttggagtlactctacgtctgagtgcatcttctcaatgggacggagcgggtgcggttccctggacagata  
20 cttctataaccaagaggagtacgtgcgttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
cctaGcgccgagtlactggaacagccagaaggacAtcctggaagacaggcgggccTggtggacacctactgcagac  
acaactacgggggttggtgagagcttcacagtgca(SAQ ID NO: 2 1 0) ;

DRB1\*131401 :

lacgtctgagtgcatcttctcaatgggacggagcgggtgcggttccctggacagatacttctataaccaagaggag  
25 tAcgtgcgttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcgccgatgccgagtlactgga  
acagccagaaggacTtcttggaaGaCaggcgggccgcggtggacacctactgcagacacaactacgggggttggtg(  
SAQ ID NO: 2 1 1) ;

DRB1\*131402 :

cacgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggttccctggacagata  
cttctataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagTccgggcgggtgacggagctggggcgg  
cctgacgcTgagtacttgaacagccagaaggacTtccctggaaGaCaggcgggccgcggtggacacctactgcagac  
5 acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 2 1 2) ;

DRB1\*1315 :

tttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggttccctggaGagatacttc  
cataaccaggaggagaacgtgcgcttcgacagcgacgtgggggagTccgggcgggtgacggagctggggcggcctg  
atgccgagtagtgaacagccagaaggacatcctggaagacGAgcgggccgcggtggacacctactgcagacacaa  
10 ctacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 1 3) ;

DRB1\*1316 :

ggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggttccctggacagatacttccataac  
caggaggagaacgtgcgcttcgacagcgacgtgggggagTccgggcgggtgacggagctggggcggcctgatgccg  
agtagtgaacagccagaaggacatcctggaagacgagcgggccgcggtggacacctactgcagacacaactacgg  
15 ggttgAtgagagcttcaca (SEQ ID NO: 2 1 4) ;

DRB1\*1317 :

ggggacaccagaccacgtttcttggagtactctacgggtgagtgTatttcttcaatgggacggagcgggtgcggt  
tccctggacagatacttctataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagTccgggcgggtgac  
ggagctggggcggcctgatgccgagtagtgaacagccagaaggacatcctggaagacGAgcgggccgcggtggac  
20 acctactgcagacacaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 1 5) ;

DRB1\*1318 :

cacgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggttccctggacagata  
cttccataaccaggaggagAacgtgcgcttcgacagcgacgtgggggagTccgggcgggtgacggagctggggcgg  
cctgatgccgagtagtgaacagccagaaggacTtccctggaagacaggcggggccTggtggacacctactgcagac  
25 acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 1 6) ;

DRB1\*1319 :

ggggacaccagaccacgtttcttggagtactctacgtctgagtgatcatttcttcaatgggacggagcgggtgcggt

tcctggaGagatacttccataaccaggaggagTcgtgcgcttcgacagcgacgtgggggagtaccgggcggtgac  
ggagctggggcgccctgatgccgagctaciggaacagccagaaggacatccctggaagacGAgcgggcccgggtggac  
acctactgcagacacaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 1 7) ;  
DRB1\*1320 :

5 cacgtttcttggagtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttccttggaCagata  
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cctgatgccgagctactggaacagccagaaggacctcttgaagacGAgcgggcccgggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcgg (SEQ ID NO: 2 1 8) ;  
DRB1\*1321 :

10 ggggacaccagaccagtttcttggagtactctacgtCtgagtgctcatttcttcaatgggacggagcgggtgcggt  
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ggagctggggcgccctacGcggcagctactggaacagccagaaggacTtccctggaagaCaggcgggcccgggtggac  
acctactgcagacacaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 1 9) ;  
DRB1\*1322 :

15 gaccacgtttcttggagtactctacgtCtgagtgctcatttcttcaatgggacggagcgggtgcggttccttggaCag  
atacttctataaccaagaggagTAcgtgcgcttcgacagcgacgtgggggagtTccgggcccgtgacggagctgggg  
cgccctgatgccgagctactggaacagccagaaggacAtccctggaagacGAgcgggcccgggtggacacctactgca  
gacacaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 2 0) ;  
DRB1\*1323 :

20 cgtttcttggagtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttccttggaCagatact  
tctataaccaagaggagctacgtgcgcttcgacagcgacgtgggggagtTccgggcccgtgacggagctggggcgcc  
tgatgccgagctactggaacagccagaaggacatccctggaagacGAgcgggcccgggtggacacctactgcagacac  
aactacggggttgtGgagagcttcacGgtgcagcggc (SEQ ID NO: 2 2 1) ;  
DRB1\*1324 :

25 cgtttcttggagtactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttccttggaCagatact  
tctataaccaagaggagctacgtgcgcttcgacagcgacgtgggggagtTccgggcccgtgacggagctggggcgcc  
tgatgccgagctactggaacagccagaaggacTtccctggaagacGAgcgggcccgggtggacacctactgcagacac

aactacggggttgtGgagagcttcacagtcagcggc (SEQ ID NO: 2 2 2) ;

DRB1\*1325 :

cacgtttcttggagttactCtaCgtCtgaGtgtcatttcttcaatgggacggagcgggtgcggttccctggaCagata  
cttctataaccaagaggagtAcgtgcgttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcgg  
5 cctgatgccgagttactggaacagccagaaggacctccctggaagaCaggcgggccggtggacacctactgcagac  
acaactacggggttggtgaga (SEQ ID NO: 2 2 3) ;

DRB1\*1326 :

cacgtttcttggagttactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttccctggagagata  
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10 cctgacgtgagttactggaacagccagaaggacttccctggaGaCaggcgCggcggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 2 2 4) ;

DRB1\*1327 :

cacgtttcttggagttactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttAcctggacagata  
cttccataaaccaggaggagaacgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
15 cctgatgccgagttactggaacagccagaaggacatccctggaagacGAgcgggccggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 2 5) ;

DRB1\*1328 :

tggagttactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttccctggacagatacttccata  
accaggaggagaacgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgatgc  
20 cgagttactggaacagccagaaggacatccctggaagacgagcgggccggtggacacctactgcagacacaactac  
Cgggttggtgagagcttcac (SEQ ID NO: 2 2 6) ;

DRB1\*1329 :

cacgtttcttggagttactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttccctggaCagata  
cttccataaaccaggaggagAACgtgcgttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcgg  
25 cctgatgccgagttactggaacagccagaaggacctccctggaagacGAgcgggccggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 2 2 7) ;

DRB1\*1330 :

tttcttggagtactctacgtctgagtgatgttcttcaatgggacggagcgggtgcggttccctggacagatacttc  
tataaccaagaggagtagtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggccta  
GgcccagtagtggaaacagccagaaggacAtcctggaagaCaggcgggccggtggacacctactgcagacacaa  
ctacgggggttggtgagagcttcaca (SEQ ID NO: 2 2 8) ;

5 DRB1\*1331 :

cacgtttcttggagtactctacgtctgagtgatgttcttcaatgggacggagcgggtgcggttccctggacagata  
cttccataaaccaggaggagaacgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
cctgTgcccagtagtggaaacagccagaaggacAtcctggaagacGAgcgggccggtggacacctactgcagac  
acaactacgggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 2 2 9) ;

10 DRB1\*1332 :

cacgtttcttggagtactctacgtctgagtgatgttcttcaatgggacggagcgggtgcggttccctggacagata  
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cctaGgcccagtagtggaaacagccagaaggacatcctggaagacGAgcgggccggtggacacctactgcagac  
acaactacgggggttggtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 3 0) ;

15 DRB1\*1333 :

cacgtttcttggagtactctacgtctgagtgatgttcttcaatgggacggagcgggtgcggttccctggacagata  
cttctataaccaagaggagtagtgcgcttcgacagcgacgtgggggagttaccgggcggtgacggagctggggcgg  
cctaGgcccagtagtggaaacagccagaaggacatcctggaagaCaagcgggccggtggacaActactgcagac  
acaactacgggggttggtg (SEQ ID NO: 2 3 1) ;

20 DRB1\*1334 :

cacgtttcttggagtactctacgtctgagtgatgttcttcaatgggacggagcgggtgcggttccctggacagata  
cttccataaaccaggaggagaacCtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
cctgatgccgagtagtggaaacagccagaaggacatcctggaagacGAgcgggccggtggacacctactgcagac  
acaactacgggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 2 3 2) ;

25 DRB1\*1335 :

cacgtttcttggagtactctacgtctgagtgatgttcttcaatgggacggagcgggtgcggttccctggacagata  
cttccataaaccaggaggagaacgtgcgcttcgacagcgacgtgggggagttccTggcggtgacggagctggggcgg

cctgatgccgagtlacttggaacagccagaaggacatccctggaagacgagcgggccggttgacacctactgcagac  
acaactacggggttggtgagagcttcacagtcagcggg(SEQ ID NO: 2 3 3) ;

DRB1\*1336 :

cacgtttcttggagtlactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttccctggaCagata  
5 cttccataaccaggaggagAACgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
cctgatgccgagtlacttggaacagccagaaggacAtccctggaagacGAgcgggccggttgacacctactgcagac  
acaactacggggttggtgagagcttcacagtcagcggcgag(SEQ ID NO: 2 3 4) ;

DRB1\*1337 :

cacgtttcttggagtlactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttccctggacagata  
10 cttctataaccaagaggagtlacgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
cctgatgccgagtlacttggaacagccagaaggacatccctggaagaCaAgcgggccggttgacacctactgcagac  
acaactacggggttggtgagagcttcacGgtgcagcggcga(SEQ ID NO: 2 3 5) ;

DRB1\*1338 :

cacgtttcttggagtlactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttccctggacagata  
15 cttctataaccaagaggagtlacgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
ccttGcgccgagtlacttggaacagccagaaggacatccctggaagacGAgcgggccggttgacacctactgcagac  
acaactacggggttggt(SEQ ID NO: 2 3 6) ;

DRB1\*1339 :

cacgtttcttggagtlactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttccctggacagata  
20 cttccataaccaggaggagaacgtgcgcttcgacagcgacgtgggggagtlccgggcggtgacggagctggggcgg  
cctgatgccgagtlacttggaacagccagaaggacAtccctggaagacGAgcgggccggttgacacctactgcagac  
acaactacggggttggtgagagcttcacagtcagcggcgag(SEQ ID NO: 2 3 7) ;

DRB1\*1340 :

ttggagtlactctacgtctgagtgctcatttcttcaatgggacggagcgggtgcggttccctggaCagatacttccata  
25 accaggaggagAACgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcggccctgatgc  
cgagtlacttggaacagccagaaggacAtccctggaagacGAgcgggccggttgacacctactgcagacacaactac  
ggggttgtGgagagcttcacagtcagcggcg(SEQ ID NO: 2 3 8) ;



DRB1\*1341 :

cacgtttcttggaglac tctacgtctgagtgat tttcttcaatgggacggagcgggtgcggtAcc tggacagata  
cttccataaaccaggaggagaacgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
cctgatgccgagtlactggaacagccagaaggacatccttgggaagacGAgcggggccgcggtggacacctactgcagac  
5 acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 2 3 9) ;

DRB1\*1342 :

cacgtttcttggaglac tctacgtctgagtgat tttcttcaatgggacggagcgggtgcggttcc tggacagata  
cttccataaaccaggaggagAacgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
cctgatgccgagtlactggaacagccagaaggacTtcttgggaagaCaggcggggccgcggtggacacctactgcagac  
10 acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 4 0) ;

DRB1\*1343 :

cacgtttcttggaglac tctacgtctgagtgat tttcttcaatgggacggagcgggtgcggttcc tggacagata  
cttccataaaccaggaggagaacgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
cctgCtgcgggagcac tggaaacagccagaaggacctccttgggaagacGAgcggggccgcggtggacacctactgcagac  
15 acaactacggggttgtGgagagcttcacagtcagcggcgga (SEQ ID NO: 2 4 1) ;

DRB1\*1344 :

cacgtttcttggaglac tctacgtCtgaGtgat tttcttcaatgggacggagcgggtgcggttcc tggacagata  
cttctataaccaagaggagtAcgtgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
cctgatgccgagtlactggaacagccagaaggacctccttggagcagaggcggggccgcggtggacacctactgcagac  
20 acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 4 2) ;

DRB1\*1345 :

t tggaglac tctacgtctgagtgat tttcttcaatgggacggagcgggtgcggttcc tggacagatacttctata  
accaagaggagtlactgcgttcgacagcgacgtgggggagttccgggcggtgacggagctggggcggcctgCtgc  
ggagcac tggaaacagccagaaggacAtccttgggaagacGAgcggggccgcggtggacacctactgcagacacaactac  
25 ggggttggtagag (SEQ ID NO: 2 4 3) ;

DRB1\*1346 :

cacgtttcttggaglac tctacgtctgagtgat tttcttcaatgggacggagcgggtgcggttcc tggacagata

cttctataaccaagaggagtAcgtgcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
ccigtTgcccgagtiactggaacagccagaaggacTtccitggaAgaCaggcgggccggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 2 4 4) ;

DRB1\*1347 :

- 5 cactttcttggagtactctacgtCtgagtgtcatitcttcaatgggacggagcgggtgcggttccitggacagata  
cttctataaccaagaggagtAcgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
cctgatgccgagtiactggaacagccagaaggacTtccitggaagacaggcggggcccTggaggacacctactgcagac  
acaactacggggttggtgagagcttcacGgtgcagcggcgag (SEQ ID NO: 2 4 5) ;

DRB1\*1348 ;

- 10 cactttcttggagtactctacgtCtgagtgtcatitcttcaatgggacggagcgggtgcggttccitggacagata  
cttctataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
cctaGgcccgagtiactggaacagccagaaggacatccitggaagacGAgcggggccggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 4 6) ;

DRB1\*1349 :

- 15 cactttcttggagtactctacgtCtgagtgtcatitcttcaatgggacggagcgggtgcggttccitggacagata  
cttctataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
cctaGgcccgagtiactggaacagccagaaggacTtccitggaagaCaggcggggccggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtcagcgg (SEQ ID NO: 2 4 7) ;

DRB1\*1350 :

- 20 cactttcttggagtactctacgtCtgagtgtcatitcttcaatgggacggagcgggtgcggttccitggacagata  
cttctataaccaagaggagAacgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
cctgatgccgagtiactggaacagccagaaggacTtccitggaagaCaggcggggccggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 2 4 8) ;

DRB1\*1351 :

- 25 cactttcttggagtactctacgtCtgagtgtcatitcttcaatgggacggagcgggtgcggttccitggacagata  
cttccataaccaggaggagaacgtgcgcttcgacagcgacgtgggggagtlccgggcgTigacggagctggggcgg  
cctgatgccgagtiactggaacagccagaaggacatccitggaagacgagcggggccggtggacacctactgcagac

acaactacggggttgttgagagcttcacagtcagcggcgag (SEQ ID NO: 2 4 9) ;

DRB1\*1352 :

cacgtttcttggagtactctacgtctgagtgctatttcttcaatgggacggagcgggtgcggttccttggaCagata  
cttccataaccaGgaggagtAcgtgcgcttcgacagcgacgtgggggagTccgggcggtagcggagctggggcgg  
5 cctgatgccgagtagtggaaacagccagaaggacAtcctggaagacGAgcgggcccggtagcacctactgcagac  
acaactacggggttgtGgagagcttcacagt (SEQ ID NO: 2 5 0) ;

DRB1\*1353 :

cacgtttcttggagtactctacgtctgagtgctatttcttcaatgggacggagcgggtgcggttccttggaGagata  
cttccataaccaggaggagaAcgtgcgcttcgacagcgacgtgggggagTaccgggcggtagcggagctggggcgg  
10 cctgatgccgagtagtggaaacagccagaaggacAtcctggaagacGAgcgggcccggtagcacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgga (SEQ ID NO: 2 5 1) ;

DRB1\*1354 :

cacgtttcttggagtactctacgtctgagtgctatttcttcaatgggacggagcgggtgcggttccttggaCagata  
cttctataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagTccgggcggtagcggagctggggcgg  
15 cctgtcgccgagTcctggaacagccagaaggacttcctggaagacGAgcgggcccggtagcacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 5 2) ;

DRB1\*1355 :

tttcttggagtactctacgtCtgagtgctatttcttcaatgggacggagcgggtgcggttccttggaCagatactt  
tataaccaagaggagtacgtgcgcttcgacagcgacgtgggggagTaccgggcggtagcggagctggggcggccta  
20 gCgccgagtagtggaaacagccagaaggacTtctggaagacaggcgggccTggtggacacctactgcagacacaa  
ctacggggttggtgagagcttcacGgtgcagcggcgag (SEQ ID NO: 2 5 3) ;

DRB1\*140101 :

atgggtgtgtctgaggctccctggaggctccgtcaltggcagTctgacagtgacactgatgggtgtgagctccccac  
tggctttggctggggacaccagaccagtttcttggagtactctacgtCtgagtgctatttcttcaatgggacgga  
25 gcgggtgcggttccttggaCagatacttccataaccaggaggagTtcgtgcgcttcgacagcgacgtgggggagTac  
cgggcggtagcggagctggggcggccctgtcgtcgagacactggaacagccagaaggaccttcctggagcggaggcggg  
ccgAggtggacacctatTgcagacacaaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID

NO: 2 5 4) ;

DRBI\*140102 :

cacgttcttggagtlacTctacgtCtgagtgtcatttcttcaatgggacggagcgggtgcggttcttggacagata  
cttccataaaccaggaggagttcgtgcgttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
5 cctgCtgcgggagcactggaacagccagaaggaccttctggagcggaggcgggcccAggtggacacctactgcagac  
acaactacggggttgtGg (SEQ ID NO: 2 5 5) ;

DRBI\*1402 :

atgggtgtgtctgaggctccctggaggctccttgcatggcagTtctgacagtgacactgatgggtgtgagctccccac  
tggctttggctggggacaccagaccagtttcttggagtlactctacgtctgagtgtcatttcttcaatgggacgga  
10 gcgggtgcggttcttggagagatacttccataaccaGaggagAAcgtgcgttcgacagcgacgtgggggagtlac  
cgggcggtgacggagctggggcggccigtatgccgagtlactggaacagccagaaggaccttctggagcagaggcggg  
ccgcggtggacacctactgcagacacaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID  
NO: 2 5 6) ;

DRBI\*1403 :

15 atgggtgtgtctgaggctccctggaggctccttgcatggcagTtctgacagtgacactgatgggtgtgagctccccac  
tggctttggctggggacaccagaccagtttcttggagtlactctacgtctgagtgtcatttcttcaatgggacgga  
gcgggtgcggttcttggagagatacttccataaaccaggaggagAACgtgcgttcgacagcgacgtgggggagtlac  
cgggcggtgacggagctggggcggccigtatgccgagtlactggaacagccagaaggaccttctggaagacaggcggg  
cccTggtggacacctactgcagacacaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID

20 NO: 2 5 7) ;

DRBI\*1404 :

atgggtgtgtctgaggctccctggaggctccttgcatggcagTtctgacagtgacactgatgggtgtgagctccccac  
tggctttggctggggacaccagaccagtttcttggagtlactctacgggtgagtgTatttcttcaatgggacgga  
gcgggtgcggttcttggacagatacttccataaaccaggaggagttcgtgcgttcgacagcgacgtgggggagtlac  
25 cgggcggtgacggagctggggcggccigtatgccgagcactggaacagccagaaggaccttctggagcggaggcggg  
ccgAggtggacacctatTgcagacacaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID  
NO: 2 5 8) ;

DRB1\*140501 :

cacgtttcttggagtaacctacgtctgagtgtaatttcttcaatgggacggagcgggtgcggttccctggacagata  
cttccataaaccaggaggagttcgtgcgttcgacagcgacgtgggggagtagcgggctggggcgg  
cctgatgcTgagtaactggaacagccagaaggacctccctggagcggaggcgggcccagggtggacacctatTgcagac  
5 acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 5 9) ;

DRB1\*140502 :

cacgtttcttggagtaacctacgtctgagtgtaatttcttcaatgggacggagcgggtgcggttccctggacagata  
cttccataaaccaggaggagttcgtgcgttcgacagcgacgtgggggagtagcgggctggggcgg  
cctgatgccgagtaactggaacagccagaaggacctccctggagcggaggcgggcccagggtggacacctatTgcagac  
10 acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 6 0) ;

DRB1\*1406 :

cacgtttcttggagtaacctacgtctgagtgtaatttcttcaatgggacggagcgggtgcggttccctggaGagata  
cttccataaaccaggaggagAACgtgcgttcgacagcgacgtgggggagtagcgggctggggcgg  
cctgatgccgagtaactggaacagccagaaggacctccctggagcagaggcgggcccgggtggacacctactTgcagac  
15 acaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 2 6 1) ;

DRB1\*140701 :

cacgtttcttggagtaacctacgtctgagtgtaatttcttcaatgggacggagcgggtgcggttccctggacagata  
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cctgCtgcggagcactggaacagccagaaggacctccctggagcggaggcgggcccAggtggacacctatTgcagac  
20 acaactacggggttgtGgagagcttcacagtcagcggcga (SEQ ID NO: 2 6 2) ;

DRB1\*140702 :

cacgtttcttggagtaacctacgtctgagtgtaatttcttcaatgggacggagcgggtgcggttccctggacagata  
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cctgCtgcggagcactggaacagccagaaggacctccctggagcggaggcgggcccagggtggacacctatTgcagac  
25 acaactacggggttgtGgagagcttcacGgtgcagcggcgag (SEQ ID NO: 2 6 3) ;

DRB1\*1408 :

cacgtttcttggagtaacctacgtctgagtgtaatttcttcaatgggacggagcgggtgcggttccctggacagata

cttccataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagtagcgggaggtagcggagctggggcgg  
cctgatgcggagCactggaacagccagaaggacctcctggagcggaggcgggccgAggtggacacctatTgcagac  
acaactacggggttgtGgagagcttcacagtgcagcggcga (SEQ ID NO: 264) ;

DRB1\*1409 :

- 5 tttcttggagtagtcttaCgtctgaGtgctatttttccaatgggacggagcgggtgcggttccctggaCagatacttc  
CataaccaGaggagAACgtgcgcttcgacagcgacgtgggggagtagcgggaggtagcggagctggggcggcctg  
atgccgagtactggaacagccagaaggacctcctggagcagaggcgggccgaggtagcggagacctactgcagacacaa  
ctacggggttgtGgagagcttcacagtgcagcggcga (SEQ ID NO: 265) ;

DRB1\*1410 :

- 10 tttcttggagcaggttaaACtagtgctatttttccaatgggacggagcgggtgcggttccctggacagatacttcc  
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tgaggagcactggaacagccagaaggacctcctggagcggaggcgggccgAggtggacacctatTgcagacacaa  
ctacggggttgtGgagagcttcacagtgcagcgg (SEQ ID NO: 266) ;

DRB1\*1411 :

- 15 gagtactctacgggtgagtgTatttttccaatgggacggagcgggtgcggttccctggacagatacttccataacc  
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gttgtGg (SEQ ID NO: 267) ;

DRB1\*1412 :

- 20 gtctgagtgctatttttccaatgggacggagcgggtgcggttccctggaGagatacttccataaccaggaggagAAC  
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ID NO: 268) ;

DRB1\*1413 :

- 25 gagtactctacgtctgagtgctatttttccaatgggacggagcgggtgcggttccctggaGagatacttccataacc  
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gtactggaacagccagaaggacctcctggagcagaggcgggccgaggtagcggagacctactgcagacacaaactacggg

gttggig(SEQ ID NO: 2 6 9) ;

DRB1\*1414 :

tcttggagtactctacgtCtgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagatacttcc  
ataaccaggaggagtTcgtgcgttcgacagcgacgtgggggagtagcgggcgggtgacggagctggggcggcctga  
5 tgcggagtactggaacagccagaaggacctccctggagcggaggcgggcccAggtggacacctatTgcagacacaa  
tacggggttggtgagagcttcacagt(SEQ ID NO: 2 7 0) ;

DRB1\*1415 :

ctctacgggtgagtgTatttcttcaatgggacggagcgggtgcggttccctggacagatacttccataaccaggag  
gagtTcgtgcgttcgacagcgacgtgggggagtagcgggcgggtgacggagctggggcggcctgaatgccgagtact  
10 ggaacagccagaaggacTtccctggaagacaggcgggcccTggtggacacctactgcagacacaaactacggggttgt  
Ggagagcttcacagtgcag(SEQ ID NO: 2 7 1) ;

DRB1\*1416 :

tiggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagatacttccata  
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15 ggagcactggaacagccagaaggacAtcctggaagacGAgcgggcccgggtggacacctactgcagacacaaactac  
ggggttgtGgag(SEQ ID NO: 2 7 2) ;

DRB1\*1417 :

cagtttcttggagtactctacgtctgaGtgatcttcttcaatgggacggagcgggtgcggttccctggaCagata  
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20 cctgatgccgagtactggaacagccagaaggacctccctggagcagaggcgggcccgggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtgcagcgg(SEQ ID NO: 2 7 3) ;

DRB1\*1418 :

gagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttccctggagagatacttccataacc  
aggaggagAACgtgcgttcgacagcgacgtgggggagtagcgggcgggtgacggagctggggcggcctgatgcTga  
25 gtactggaacagccagaaggacctccctggagcggaggcgggcccAggtggacacctatTgcagacacaaactacggg  
gttgtGgagagcttcacagtgcagcggcga(SEQ ID NO: 2 7 4) ;

DRB1\*1419 :

ggggacaccagaccacgtttcttggAgtaactcttaCgtctgagtgtaatttcttcaatgggacggagcgggtgcggt  
tcttggagagatacttccataaccaggaggagAACgtgcgttcgacagcgacgtgggggagtagcgggcgggtgac  
ggagctggggcggcctgatgccgagtagtgaacagccagaaggacctcctggagcagaAgcgggccgcggtggac  
acctactgcagacacaactacggggttggtgagagcttcaca (SEQ ID NO: 275) ;

## 5 DRB1\*1420 :

tggagtagtcttacgtCtgagtgtaatttcttcaatgggacggagcgggtgcggttCcttggagagatacttccata  
accaggaggagTcgtgcgttcgacagcgacgtgggggagtagcgggcgggtgacggagctggggcggcctgatgc  
cgagtagtgaacagccagaaggacctcctggagcagaggcgggccgcggtggacacctactgcagacacaactac  
ggggttgtGgaga (SEQ ID NO: 276) ;

## 10 DRB1\*1421 :

tggagtagtcttacgtCtgagtgtaatttcttcaatgggacggagcgggtgcggttCcttggagagatacttccata  
accaggaggagAACgtgcgttcgacagcgacgtgggggagtagcgggcgggtgacggagctggggcggcctgatgc  
cgagtagtgaacagccagaaggacctcctggagcagaAgcgggccgcggtggacacctactgcagacacaactac  
ggggttgtGgaga (SEQ ID NO: 277) ;

## 15 DRB1\*1422 :

cacgtttcttggagtagtcttacgtCtgagtgtaatttcttcaatgggacggagcgggtgcggttCcttggagagata  
cttccataaccaggaggagTcgtgcgttcgacagcgacgtgggggagtagcgggcgggtgacggagctggggcgg  
cctgCtgcggagCactgaacagccagaaggacTtcttggagagaCaggcgggccgcggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtgcagcggcgag (SEQ ID NO: 278) ;

## 20 DRB1\*1423 :

cacgtttcttggagtagtcttacgtCtgagtgtaatttcttcaatgggacggagcgggtgcggttCcttggagagata  
cttccataaccaggaggagTcgtgcgttcgacagcgacgtgggggagtagcgggcgggtgacggagctggggcgg  
cctgatgccgagtagtgaacagccagaaggacctcctggagcggaggcgggccgAggtggacacctatTgcagac  
acaactacggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 279) ;

## 25 DRB1\*1424 :

tcttggagtagtcttacgtCtgagtgtaatttcttcaatgggacggagcgggtgcggttCcttggagagatacttcc  
ataaccaGgaggagAACgtgcgttcgacagcgacgtgggggagtagcgggcgggtgacggagctggggcggcctga



tgccgagtacttggaacagccagaaggacAtcctggagcagGCgcgggccggtggacacctactgcagacacaa  
tacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 280) ;

DRB1\*1425 :

ttcttggagtactctacgtCtgagtgctatttcttcaatgggacggagcgggtgcgggttccctggacagatacttc  
5 tataaccaagaggagAACgtgcgttcgacagcgacgtgggggagtlaccgggcgggtgacggagctggggcgccctg  
CtgcggagCacttggaacagccagaaggacTtccctggaAgaCaggcgggccggtggacacctactgcagacacaa  
ctacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 281) ;

DRB1\*1426 :

cacgtttcttggagtactctacgtctgagtgctatttcttcaatgggacggagcgggtgcAgttccctggacagata  
10 ctccataaaccaggaggagttcgtgcgttcgacagcgacgtgggggagtlaccgggcgggtgacggagctggggcg  
cctgctgcgggagcacttggaacagccagaaggacctccctggagcggaggcgggccgaggtggacacctattgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 282) ;

DRB1\*1427 :

cacgtttcttggagtactctacgtctgagtgctatttcttcaatgggacggagcgggtgcgggttccctggaGagata  
15 ctccataaaccaggaggagAACgtgcgttcgacagcgacgtgggggagtlaccgggcgggtgacggagctggggcg  
cctgatgccgagtacttggaacagccagaaggacTtccctggaagaCaggcgggccCTggtggacacctactgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 283) ;

DRB1\*1428 :

cacgtttcttggagtactctacgggtgagtgctatttcttcaatgggacggagcgggtgcgggttccctggacagata  
20 ctccataaaccaggaggagttcgtgcgttcgacagcgacgtgggggagtlaccgggcgggtgacggagctggggcg  
cctgCtgcgggagcacttggaacagccagaaggacctccctggagcggaggcgggccgaggtggacacctattgcagac  
acaactacggggCtgtGgagagcttcaca (SEQ ID NO: 284) ;

DRB1\*1429 :

cacgtttcttggagtactctacgtctgagtgctatttcttcaatgggacggagcgggtgcgggttccctggaGagata  
25 ctccataaccaGaggagAACgtgcgttcgacagcgacgtgggggagtlaccgggcgggtgacggagctggggcg  
cctgatgccgagtacttggaacagccagaaggacctccctggagcagaggcgggccggtggacacctactgcagac  
acaactacggggCtgtggagagcttcacagtcagcggcgag (SEQ ID NO: 285) ;

DRB1\*1430 :

tttcttggagtactctacgtctgaGtgtcattttcttcaatgggacggagcgggtgcggttccctggaCagatacttc  
CataaccaGgaggagAacgtgcgcttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcggcctg  
atgccgagtactggaacagccagaaggaccttctggagcagaggcgggccggtggacacctactgcagacacaa  
5 ctacgggggttggtagagcttcaca (SEQ ID NO: 286) ;

DRB1\*1431 :

tttcttggagtactctacgggtgagtgTattttcttcaatgggacggagcgggtgcggttccctggacagatacttc  
cataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagttaccgggcggtgacggagctggggcggcctg  
CtgcgggagcactggaacagccagaaggaccttctggagcggaggcgggccggtggacacctatTgcagacacaa  
10 ctacgggggttgtGgagagcttcaca (SEQ ID NO: 287) ;

DRB1\*1432 :

cacgtttcttggagtacTctacgtCtgagtgtcattttcttcaatgggacggagcgggtgcggttccctggacagata  
cttccataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
ccctgCtgcgggagcactggaacagccagaaggaccttctggagcGgaggcgggccggtggacacctactgcagac  
15 acaactacgggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 288) ;

DRB1\*1433 :

tiggagtactctacgtctgagtgTcattttcttcaatgggacggagcgggtgcggttccctggacagatacttccata  
accaggaggagaAcgtgcgcttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcggcctgatgc  
cgagtactggaacagccagaaggaccttctggagcagaggcgggccgAggtggacacctactgcagacacaactac  
20 ggggttgtGgagagcttcacagtgcagcggc (SEQ ID NO: 289) ;

DRB1\*1434 :

cacgtttcttggagtacTctacgtCtgagtgtcattttcttcaatgggacggagcgggtgcggttccctggacagata  
cttccataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
ccctgatgcggagCactggaacagccagaaggaccttctggagcggaggcgggccggtggacacctatTgcagac  
25 acaactacgggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 290) ;

DRB1\*1435 :

cacgtttcttggagtactctacgtctgagtgTcattttcttcaatgggacggagcgggtgcggttccctggacagata

cttccataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagtTccgggcggtgacggagctggggcgg  
ccigCtgcgggagcactggaacagccagaaggacctcctggagcggaggcgggcccAggtggacacctatTgcagac  
acaactacggggttgtGgagagcttcacagtcagcggg(SEQ ID NO: 2 9 1) ;

DRB1\*1436 :

- 5 cacttttcttggagtactctacgtctgagtgicatttcttcaatgggacggagcgggtgcggttccitggacagata  
cttccataaccaggaggagttcgtgcgcttcgacagcgacgtGgggagtlaccgggcggtgacggagctggggcgg  
ccigtatgccgagtactggaacagccagaaggacctcctggagcggaggcgggcccagggtggacacctatTgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag(SEQ ID NO: 2 9 2) ;

DRB1\*1437 :

- 10 cacttttcttggagtactctacgtctgagtgicatttcttcaatgggacggagcgggtgcggttccitggacagata  
cttccataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
ccigtatgctgagtactggaacagccagaaggacctcctggagcaggCgcgggcccgggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag(SEQ ID NO: 2 9 3) ;

DRB1\*1438 :

- 15 cacttttcttggagtactctacgtctgagtgicatttcttcaatgggacggagcgggtgcggttccitggacagata  
cttccataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
ccigCtgcgggagcactggaacagccagaaggacctcctggagcggaggcgggcccagggtggacaaTtactgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag(SEQ ID NO: 2 9 4) ;

DRB1\*1439 :

- 20 cacttttcttggagtacctacgtCtgagtgicatttcttcaatgggacggagcgggtgcggttccitggacagata  
cttccataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
ccigCtgcgggagcactggaacagccagaaggacctcctggagcggaggcgggcccAggtggacacctatTgcagac  
acaactacggggttgtGgagagcttcacagtcagcggcgag(SEQ ID NO: 2 9 5) ;

DRB1\*1440 :

- 25 cacttttcttggagtactctacgtctgagtgicatttcttcaatgggacggagcgggtgcggttccitggaGagata  
cttccataaccaggaggagtTcgtgcgcttcgacagcgacgtgggggagtlaccgggcggtgacggagctggggcgg  
ccigtatgccgagtactggaacagccagaaggacctcctggaagaCaggcgggcccTggtggacacctactgcagac

acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 296) ;

DRB1\*1441 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttcttggaGagata  
cttccataaaccaggaggagtTcTtgcgcttcgacagcgacgtgggggagtagcgggctggggcgg

5 cctgatgccgagtacttggaacagccagaaggacctcttgagcagaggcgggcccgggtggacacctactgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 297) ;

DRB1\*1442 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttcttggaGagata  
cttctataaccaagaggagtAcgtgcgcttcgacagcgacgtgggggagtagcgggctggggcgg

10 cctgatgccgagtacttggaacagccagaaggacctcttgagcggaggcgggcccAggtggacacctatTgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 298) ;

DRB1\*1443 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttcttggaGagata  
cttccataaaccaggaggagttcgtgcgcttcgacagcgacgtgggggagtagcgggctggggcgg

15 cctgatgctgagtacttggaacagccagaaggacctcttgagcggaggcgggcccagggtggacGcctatTgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 299) ;

DRB1\*1444 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttcttggaGagata  
cttccataaaccaggaggagttcgtgcgcttcgacagcgacgtgggggagtagcgggctggggcgg

20 cctgatgctgagtacttggaacagccagaaggacctcttgagcggaggcgggcccagggtggacacctatTgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 300) ;

DRB1\*1445 :

cacgtttcttggagtactctacgtctgagtgatcttcttcaatgggacggagcgggtgcggttcttggaGagata  
cttccataaaccaggaggagttcgtgcgcttcgacagcgacgtgggggagtagcgggctggggcgg

25 cctgatgctgagtacttggaacagccagaaggacctcttgagcggaggcgggcccagggtggacacctatTgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 301) ;

DRB1\*150101 :

atggtgtgtctgaagctccctggaggctccctgcatgacagcgctgacagtgacactgaagggtgtgagctccccac  
tggcttttgTctlggggacacccgaccacgtttccctgtggcagccctaaagaggagtgatcttcttcaatgggacgga  
gcgggtgcggttccctggacagatacttctataaccaggaggagtcctgtcgcttcgacagcgacgtgggggagttc  
cgggcggtgacggagctggggcggcctgacgctgagtactggaacagccagaaggacatccctggagcaggcgcggg  
5 ccgcggtggacacctactgcagacacaactacggggttgtggagagcttcacagtgcagcggcgag (SEQ ID  
NO: 3 0 2) ;

DRB1\*150102 :

cacgtttccctgtggcagccctaaagaggagtgatcttcttcaatgggacggagcgggtgcggttccctggacagata  
cttctataaccaggaggagtcctgtcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
10 cctgacgctgagtactggaacagccagaaggacatccctggagcaggcgcgggccgcggtggacacctactgcagac  
acaactacggAgttgtGgagagcttcacagtgcagcgg (SEQ ID NO: 3 0 3) ;

DRB1\*150103 :

cacgtttccctgtggcagccctaaagGgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagata  
cttctataaccaggaggagtcctgtcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
15 cctgacgctgagtactggaacagccagaaggacatccctggagcaggcgcgggccgcggtggacacctatTgcagac  
acaactacggggttgtGgagagcttcacagtgcagcgg (SEQ ID NO: 3 0 4) ;

DRB1\*150104 :

cacgtttccctgtggcagccctaaagGgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagata  
cttctataaccaggaggagtcctgtcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
20 cctgatgccgagtactggaacagccagaaggacAtcctggagcaggCgcgggcccgcggtggacacctactgcagac  
acaactacggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 3 0 5) ;

DRB1\*150201 :

ggggacacccgaccacgtttccctgtggcagccctaaagGgagtgatcttcttcaatgggacggagcgggtgcggt  
tccctggacagatacttctataaccaggaggagtcctgtcgcttcgacagcgacgtgggggagttccgggcggtgac  
25 ggagctggggcggcctgacgctgagtactggaacagccagaaggacAtcctggagcaggCgcgggcccgcggtggac  
acctactgcagacacaactacggggttgtgagagcttcacagtgcagcggcgag (SEQ ID NO: 3 0 6) ;

DRB1\*150202 :

gagtgtcatittcttcaatgggacggagcgggtgcggttccctggacagatacttctataaccaggaggagtccgtgc  
gccttcgacagcgacgtgggggagtTccgggcgggtgacggagctggggcggcctgaigccgagttacttggaacagcca  
gaaggacAtccctggagcagGcggggccgcgggtggacacctactgcagacacaactacggggttggtg (SEQ ID  
NO: 307) ;

5 DRB1\*150203 :

cacgtttccctgtggcagcctaagaggagtgcatittcttcaatgggacggagcgggtgcggttccctggacagata  
cttctataaTcaggaggagtccgtgcgcttcgacagcgacgtgggggagtTccgggcgggtgacggagctggggcgg  
cctgacgctgagttacttggaacagccagaaggacatccctggagcaggcggggccgcgggtggacacctactgcagac  
acaactacggggttggtg (SEQ ID NO: 308) ;

10 DRB1\*1503 :

ggggacacccgaccacgtttccctgtggcagcctaagagGgagtgtcatittcttcaatgggacggagcgggtgcggt  
tccctggacagaCacttctataaccaggaggagtccgtgcgcttcgacagcgacgtgggggagtTccgggcgggtgac  
ggagctggggcggcctgacgcTgagtacttggaacagccagaaggacAtccctggagcaggCgcgggccgcgggtggac  
acctactgcagacacaactacggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 309) ;

15 DRB1\*1504 :

tccctgtggcagcctaagagGgagtgtcatittcttcaatgggacggagcgggtgcggttccctggacagatacttct  
ataaccaggaggagtccgtgcgcttcgacagcgacgtgggggagtTccgggcgggtgacggagctggggcggcctga  
cgctgagttacttggaacagccagaaggacTccctggagcaggCgcgggccgcgggtggacacctactgcagacacaac  
tacggggttgtGgagagcttcacagtgcagcgg (SEQ ID NO: 310) ;

20 DRB1\*1505 :

tccctgtggcagcctaagagGgagtgtcatittcttcaatgggacggagcgggtgcggttccctggacagatacttct  
ataaccaggaggagtccgtgcgcttcgacagcgacgtgggggagtTccgggcgggtgacggagctggggcggcctga  
cgctgagttacttggaacagccagaaggacctccctggagcaggCgcgggccgcgggtggacacctactgcagacacaac  
tacggggttgtGgagagcttcacagtgcagcgg (SEQ ID NO: 311) ;

25 DRB1\*1506 :

ctgtggcagcctaagaggagtgcatittcttcaatgggacggagcgggtgcggttccctggacagatacttctata  
accaggaggagtccgtgcgcttcgacagcgacgtgggggagtTccgggcggCgacggagctggggcggcctgacgc

tgaglac tggacagccagaaggacatcctggagcaggcgccggcggtggacacctactgcagacacaactac  
gggggtgtggagagcttcacagtgcagcggcgag (SEQ ID NO: 3 1 2) ;

DRB1\*1507 :

tttccctgtggcagcc taagagGgagtgatcatttcttcaatgggacggagcgggtgcggttcc tggacagatacttc  
5 tataaccaggaggagtcctgtgcgttcgacagcgacgtgggggagtlaccggcggtgacggagctggggcgccctg  
acgcTgagtlacttggacagccagaaggacAtcctggagcaggCgcgggccgcggtggacacctactgcagacacaa  
ctacggggttgtGgagagc (SEQ ID NO: 3 1 3) ;

DRB1\*1508 :

cacgtttccctgtggcagcc taagagggagtgatcatttcttcaatgggacggagcgggtgcggttcc tggacagata  
10 cttctataaccaggaggagtcctgtgcgttcgacagcgacgtgggggagtlccggcggtgacggagctggggcg  
cc tgcgtgagtlacttggacagccagaagAacatcctggagcaggcgccggccgcggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtgcagcggcgag (SEQ ID NO: 3 1 4) ;

DRB1\*1509 :

cacgtttccctgtggcagcc taagagggagtgatcatttcttcaatgggacggagcgggtgcggttcc tggacagata  
15 cttctataaccaggaggagtcctgtgcgttcgacagcgacgtgggggagtlccAggcggtgacggagctggggcg  
cc tgcgtgagtlacttggacagccagaaggacatcctggagcaggCgcgggccgcggtggacacctactgcagac  
acaactacggggttgtggagagcttcacagtgcagcggcgag (SEQ ID NO: 3 1 5) ;

DRB1\*1510 :

gtttccctgtggcagcc taagagGgagtgatcatttcttcaatgggacggagcgggtgcggttcc tggacagatactt  
20 ctataaccaggaggagtcctgtgcgttcgacagcgacgtgggggagtlccggcggtgacggagctggggcgcc  
gacgtgagtlacttggacagccagaaggacatcctggaagacgAgcgggccgcggtggacacctactgcagacaca  
actacggggttgtGgagagc (SEQ ID NO: 3 1 6) ;

DRB1\*1511 :

cacgtttccctgtggcagcc taagagGgagtgatcatttcttcaatgggacggagcgggtgcggttcc tggacagata  
25 cttctataaccaggaggagtcctgtgcgttcgacagcgacgtgggggagtlaccggcggtgacggagctggggcg  
cc tgcgtgagtlacttggacagccagaaggacAtcctggagcaggCgcgggccgcggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtgcagcggcgag (SEQ ID NO: 3 1 7) ;

DRB1\*1512 :

gcacgtttccctgtggcagcctaagagGgagtgatcattttctcaatgggacggagcgggtgcggttccctggacagat  
acttctataaaccaggaggagtcctgtcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcg  
gcctaGCccgagttactggaacagccagaaggacAtcctggagcaggCgcgggccgcggtggacacctactgcaga  
5 cacaactacggggttgtGgagagcttcacagtcagcggcgag (SEQ ID NO: 318) ;

DRB1\*1513 :

cacgtttccctgtggcagcctaagagGgagtgatcattttctcaatgggacggagcgggtgcggttccctggacagata  
cttctataaaccaggaggagtcctgtcgcttcgacagcgacgtgggggagttccgggcggtgacggagctggggcgg  
cctgacgcTgagtactggaacagcca...ggacAtcctggagcaggCgcgggccgcggtggacacctactgcagac  
10 acaactacggggttgtGgagagcttcacagtcagcgg (SEQ ID NO: 319) ;

DRB1\*160101 :

atgggtgtgtctgaagctccctggaggctcctgcatgacagcgctgacagtgacactgatgggtgtgagctccccac  
tggctttggctggggacacccgaccacgtttccctgtggcagcctaagagGgagtgatcattttctcaatgggacgga  
gcgggtgcggttccctggacagatacttctataaaccaggaggagtcctgtcgcttcgacagcgacgtgggggagtac  
15 cgggcggtgacggagctggggcggcctgacgcctgagttactggaacagccagaaggacTtccctggaagacaggcgCg  
ccgcggtggacacctactgcagacacaaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID  
NO: 320) ;

DRB1\*160102 :

cgtttccctgtggcagcctaagagGgagtgatcattttctcaatgggacggagcgggtgcggttccctggacagatact  
20 tctataaaccaggaggagtcctgtcgcttcgacagcgacgtgggggagttaccgggcggtgacggagctggggcggcc  
tgacgcctgagtactggaacagccagaaggacTtccctggaagaCaggcgggccgcggtggacacctactgcagacac  
aaactacggggttggtgagagcttcaca (SEQ ID NO: 321) ;

DRB1\*160201 :

atgggtgtgtctgaagctccctggaggctcctgcatgacagcgctgacagtgacactgatgggtgtgagctccccac  
25 tggctttggctggggacacccgaccacgtttccctgtggcagcctaagagGgagtgatcattttctcaatgggacgga  
gcgggtgcggttccctggacagatacttctataaaccaggaggagtcctgtcgcttcgacagcgacgtgggggagttac  
cgggcggtgacggagctggggcggcctgacgcctgagttactggaacagccagaaggacctccctggaagacaggcgCg



ccgcgggtggacacctactgcagacacaactacggggttggtagagcctcacagtgcagcggcgag (SEQ ID  
NO: 3 2 2) ;

DRB1\*160202 :

tttctgtggcagcctaagagGgagtgtcatitcttcaatgggacggagcgggtgcgggttctggacagatacttc  
5 tataaccaggaggagtcctgtcgcttcgacagcgacgtgggggaglacgggcgggtgacggagctggggcggcctg  
acgctgagtlacttggaacagccagaaggacctcttgaagaCaggcgggcccgggtggacacctactgcagacacaa  
ctacggggttgggtg (SEQ ID NO: 3 2 3) ;

DRB1\*1603 :

atgggtgtgtctgaagctcccggaggcctctgcatgacagcgctgacagtgacactgaatgggtgtgagctccccac  
10 tggcctttggctggggacacccgaccacgtttctgtggcagcctaagaggagtgatitcttcaatgggacgga  
gcgggtgcgggttctggacagatacttctataaccaggaggagtcctgtcgcttcgacagcgacgtgggggagtlac  
cgggcgggtgacggagctggggcggcctgacgctgagtlacttggaacagccagaaggaccttcttggaagacaggcCg  
ccgcgggtggacacctactgcagacacaactacggggttggtagagcctcacagtgcagcggcgag (SEQ ID  
NO: 3 2 4) ;

15 DRB1\*1604 :

tggcagcctaagaggagtgatitcttcaatgggacggagcgggtgcgggttctggacagatacttctataacc  
aGgaggagtcctgtcgcttcgacagcgacgtgggggagtlaccgggcgggtgacggagctggggcggcctgacgcTga  
gtacttggaacagccagaaggacTtcttggaagaCaggcgggcccCTgggtggacacctactgcagacacaactacggg  
gttgggtg (SEQ ID NO: 3 2 5) ;

20 DRB1\*1605 :

ctgtggcagcctaagagGgagtgtcatitcttcaatgggacggagcgggtgcgggttctggacagatacttctata  
accaggaggagtcctgtcgcttcgacagcgacgtgggggagtlaccgggcgggtgacggagctggggcggcctgacgc  
tgagtlacttggaacagccagaaggacAtcttggaagacaggcgcGccgcgggtggacacctactgcagacacaactac  
ggggttgggtgag (SEQ ID NO: 3 2 6) ;

25 DRB1\*1607 :

cacgtttctgtggcagcctaagaggagtgatitcttcaatgggacggagcgggtgcgggttccCggacagata  
cttctataaccaggaggagtcctgtcgcttcgacagcgacgtgggggagtlaccgggcgggtgacggagctggggcgg

ccctgacgtgagctaciggaacagccagaaggacatccctggaagacaggcgcccggtggacacctacigcagac  
acaactacggggttggtagagcttcacagtga (SEQ ID NO: 3 2 7) ;

DRB1\*1608 :

cacgttccctgtggcagcctaagagGgagtgatcttcttcaatgggacggagcgggtgcggttccctggacagata  
5 cttctataaccaggaggagaAcgtgcgcttcgacagcgacgtgggggagtlaccggcggtgacggagctggggcgg  
ccctgacgtgagctaciggaacagccagaaggacTtccctggaagacaggcgCgccgggtggacacctacigcagac  
acaactacggggttggtagagcttcacagtgcagcggcgag (SEQ ID NO: 3 2 8) ;

DRB3\*010101 :

ggggacacccgaccacgtttcttggagctgcGtaagctgagtgatcttcttcaatgggacggagcgggtgcggt  
10 acctggaCagatacttccataaccaggaggagttcCtgcgcttcgacagcgacgtgggggagtlaccggcggtgac  
ggagctggggcggcctgtCgccgagttcctggaacagccagaaggacctccctggagcagaagcggggccGggtggac  
aaTtactgcagacacaactacggggttggtagagcttcacagtgcagcggcgag (SEQ ID NO: 3 2 9) ;

DRB3\*01010201 :

atgggtgtgtctgaagctccctggaggctccagcttggcagcgttgacagtgacacitgaatgggtgtgagctcccgac  
15 tggctttCgcctgggacacccgaccacgtttcttggagctgcGtaagctgagtgatcttcttcaatgggacgga  
gcgggtgcggtacctggacagatacttccataaccaggaggagttcctgcgcttcgacagcgacgtgggggagtlac  
cgggcgggtgacggagctggggcggcctgtcggcgagttcctggaacagccagaaggacctccctggagcagaagcggg  
gccGggtggacaattactgcagacacaactacggggttggtagagcttcacagtgcagcggcgag (SEQ ID  
NO: 3 3 0) ;

20 DRB3\*010103 :

ggggacacccgaccacgtttcttggagctgcGtaagctgagtgatcttcttcaatgggacggagcgggtgcggt  
acctggaCagatacttccataaccaggaggagttcCtgcgcttcgacagcgacgtgggggagtlaccggcggtgac  
ggagctggggcggcctgttggcgagttcctggaacagccagaaggacctccctggagcagaagcggggccGggtggac  
aaTtactgcagacacaactacggggttggtagagc (SEQ ID NO: 3 3 1) ;

25 DRB3\*010104 :

cacgttcttggagctgcGtaagctgagtgatcttcttcaatgggacggagcgggtgcggtacctggacagata  
cttccataaccaggaggagttcctgcgcttcgacagcgacgtgggggagtlaccggcggtgacggagctggggcgg

ccigtgccgagtccttggaacagccagaaggacctcctggagcagaagcggggccgggtggacaaTtactgcagac  
acaactacggAgttggig(SEQ ID NO: 3 3 2) ;

DRB3\*0102 :

ggggacacccgaccacgtttcttgagctgTGtaagctgagtgicatttcttcaatgggacggagcgggtgcggt  
5 accitggaCagatacttccataaccaggaggagttcCtgcgcttcgacagcgacgtgggggagtagcgggcgggtgac  
ggagctggggcgccctgtGccgagtccttggaacagccagaaggacctcctggagcagaagcggggccGggtggac  
aaTtactgcagacacaactacggggttggtgagagc(SEQ ID NO: 3 3 3) ;

DRB3\*0103 :

cacgtttcttgagctgcGtaagctgagtgicatttcttcaatgggacggagcgggtgcggtaccitggaGagata  
10 ctccataaccaggaggagttcCtgcgcttcgacagcgacgtgggggagtagcgggcggtagcggagctggggcgg  
ccigtGccgagtccttggaacagccagaaggacctcctggagcagaagcggggccGggtggacaaTtactgcagac  
acaactacggggttggtgagagc(SEQ ID NO: 3 3 4) ;

DRB3\*0104 :

cacgtttctcggagctgcGtaagctgagtgicatttcttcaatgggacggagcgggtgcggtaccitggaCagata  
15 ctccataaccaggaggagttcCtgcgcttcgacagcgacgtgggggagtagcgggcggtagcggagctggggcgg  
ccigtGccgagtccttggaacagccagaaggacctcctggagcagaagcggggccGggtggacaaTtactgcagac  
acaactacggggttggtgagagcttcaca(SEQ ID NO: 3 3 5) ;

DRB3\*0105 :

cacgtttcttgagctgcGtaagctgagtgicatttcttcaatgggacggagcgggtgcggtaccitgAacagata  
20 ctccataaccaggaggagttcctgcgcttcgacagcgacgtgggggagtagcgggcggtagcggagctggggcgg  
ccigtgccgagtccttggaacagccagaaggacctcctggagcagaagcggggccgggtggacaaTtactgcagac  
acaactacggggttggtgagagcttcacagtcagcggcg(SEQ ID NO: 3 3 6) ;

DRB3\*0106 :

cacgtttcttgagctgcGtaagctgagtgicatttcttcaatgggacggagcgggtgcggtaccitggaCagata  
25 ctccataaccaggaggagttcgtgcgcttcgacagcgacgtgggggagtagcgggcggtagcggagctggggcgg  
ccigtGccgagtccttggaacagccagaaggacctcctggagcagaagcggggccGggtggacaaTtactgcagac  
acaactacggggttggtgagagcttcacagtcagcggcg(SEQ ID NO: 3 3 7) ;

DRB3\*0107 :

cacgtttcttggagctgcGtaagctgagtgcatittcttcaatgggacggagcgggtgcggtacctggacagata  
cttccataaccaggaggagtlacgcgcgttcgacagcgacgtgggggagtlaccgggcggtagggagctggggcgg  
cctgatgccgagtlactggaacagccagaaggacctcctggagcagaAgcggggccAggtggacaaTtactgcagac  
5 acaactacggggttggtag(SAQ ID NO: 3 3 8) ;

DRB3\*0108 :

cacgtttcttggagctgcGtaagctgagtgcatittcttcaatgggacggagcgggtgcggtacctggacagata  
cttccataaccaggaggagAACgtgcgcgttcgacagcgacgtgggggagtlaccgggcggtagggagctggggcgg  
cctgtGcccgagtccttgaacagccagaaggacctcctggagcagaagcggggccGggtggacaaTtactgcagac  
10 acaactacggggttggtagagagcttcacagtcagcgg(SAQ ID NO: 3 3 9) ;

DRB3\*0109 :

cacgtttcttggagctgcGtaagctgagtgcatittcttcaatgggacggagcgggtgcggtacctggagagaca  
cttccataaccaggaggagtlacGcgcttcgacagcgacgtgggggagtlaccgggcggtagggagctggggcgg  
cctgtGcccgagtccttgaacagccagaaggacctcctggagcagaagcggggccGggtggacaaTtactgcagac  
15 acaactacggggttggtagagagcttcacagtcagcgg(SAQ ID NO: 3 4 0) ;

DRB3\*0110 :

cacgtttcttggagctgcGtaagctgagtgcatittcttcaatgggacggagcgggtgcggtacctggacagata  
cttccataaccaggaggagttccttAgcttcgacagcgacgtgggggagtlaccgggcggtagggagctggggcgg  
cctgtGcccgagtccttgaacagccagaaggacctcctggagcagaagcggggccgggtggacaaTtactgcagac  
20 acaactacggggttggtag(SAQ ID NO: 3 4 1) ;

DRB3\*0201 :

atgggtgtctgaagctccctggaggctccagcttggcagcgttgacagtgacactgatgggtgtgagctcccgac  
tggctttGcttggggacacccgaccacgtttcttggagctgcttaagctgagtgcatittcttcaatgggacgga  
gcgggtgcggttccctggagagacacttccataaccaggaggagtlacgcgcgttcgacagcgacgtgggggagtlac  
25 cgggcggtagggagctggggcgccctgatgccgagtlactggaacagccagaaggacctcctggagcagaagcggg  
gccaggtggacaaTtactgcagacacaaactacggggttgtGgagagcttcacagtcagcggcgag(SAQ ID  
NO: 3 4 2) ;

DRB3\*020201 :

ggggacacccgaccacgtttcttgGagctgcttaagtcigagtgtcatttcttcaatgggacggagcgggtgcggt  
tcttgagagaCacttccataaccaggaggagtagCgcgcttcgacagcgacgtgggggagtagcgggcggtgaG  
ggagctggggcggcctgatgccgagtagtgaacagccagaaggacctcttgagcagaagcggggccaggtaggac  
5 aaTtactgcagacacaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 3 4 3) ;

DRB3\*020202 :

cacgtttcttgagctgcttaagtcigagtgtcatttcttcaatgggacggagcgggtgcggttcttgagagaCa  
cttccataaccaggaggagtagCgcgcttcgacagcgacgtgggggagtagcgggcggtgaGggagctggggcgg  
cctgatgccgagtagtgaacagccagaaggacctcttgagcagaagcggggccAggtggacaActtactgcagac  
10 acaactacggggttggtg (SEQ ID NO: 3 4 4) ;

DRB3\*020203 :

cacgtttcttgagctgcttaagtcigagtgtcatttcttcaatgggacggagcgggtgcggttcttgagagaca  
cttccataaccaggaggagtagCgcgcttcgacagcgacgtgggggagtagcgggcggtgagggagctggggcgg  
cctgatgccgagtagtgaacagccagaaggacctcttgagcagaagcggggccaggtaggacaattactgcagGc  
15 acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 3 4 5) ;

DRB3\*020204 :

cacgtttcttgagctgcttaagtcigagtgtcatttcttcaatgggacggagcgggtgcggttcttgagagaca  
cttccataaccaggaggagtagCgcgcttcgacagcgacgtgggggagtagcgggcggtgaGggagctggggcgg  
cctgatgcGgagtagtgaacagccagaaggacctcttgagcagaagcggggccaggtaggacaaTtactgcagac  
20 acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 3 4 6) ;

DRB3\*0203 :

tggagctgcttaagtcigagtgtcatttcttcaatgggacggagcgggtgcggttcttgagagaCacttccata  
accaGgaggagtagcgtgcgcttcgacagcgacgtgggggagtagcgggcggtgaGggagctggggcggcctgatgc  
cgagtagtgaacagccagaaggacctcttgagcagaagcggggccaggtaggacaaTtactgcagacacaactac  
25 ggggttggtgaga (SEQ ID NO: 3 4 7) ;

DRB3\*0204 :

cacgtttcttgagctgcttaagtcigagtgtcatttcttcaatgggacggagcgggtgcggttcttgagagaca

cttccataaccaggaggagtagcgcgcttcgacagcgacgtgggggagtagcgggcggtagaggagctggggcgg  
ccgatgccgagtagtggaacagccagaaggacctcctggagcagaagcggggccGggtggacaActactgcagac  
acaactacggggttgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 3 4 8) ;

DRB3\*0205 :

- 5 cgtttcttggagctgcttaagtcgagtgatcttcttcaatgggacggagcgggtgcggttccctggaGagatact  
ttccataaccaggaggagtagCgcgcttcgacagcgacgtgggggagtagcgggcggtagaggagctggggcggc  
tgatgccgagtagtggaacagccagaaggacctcctggagcagaagcggggccaggtaggacaaTtactgcagacac  
aactacggggttggtgagagcttcacagtgcag (SEQ ID NO: 3 4 9) ;

DRB3\*0206 :

- 10 cactttcttggagctgcttaagtcgagtgatcttcttcaatgggacggagcgggtgcggttccctggagagaca  
cttccataaccaggaggagAacgCgcgcttcgacagcgacgtgggggagtagcgggcggtagaggagctggggcgg  
ccgatgccgagtagtggaacagccagaaggacctcctggagcagaagcggggccaggtaggacaaTtactgcagac  
acaactacggggttggtg (SEQ ID NO: 3 5 0) ;

DRB3\*0207 :

- 15 ttggagctgcttaagtcgagtgatcttcttcaatgggacggagcgggtgcggttccctggagagacacttccata  
accaggaggagtagcgcgcttcgacagcgacgtgggggagtagcgggcggtagaggagctggggcggcctgTCgc  
cgagtagtggaacagccagaaggacctcctggagcagaagcggggccaggtaggacaaTtactgcagacacaactac  
ggggttggtgagag (SEQ ID NO: 3 5 1) ;

DRB3\*0208 :

- 20 cactttcttggagctgcttaagtcgagtgatcttcttcaatgggacggagcgggtgcggttccctggagagaca  
cttccataaccaggaggagtagcgcgcttcgacagcgacgtgggggagtagcgggcggtagaggagctggggcgg  
cctaGCgccgagtagtggaacagccagaaggacctcctggagcagaagcggggccaggtaggacaaTtactgcagac  
acaactacggggttggtg (SEQ ID NO: 3 5 2) ;

DRB3\*0209 :

- 25 cactttcttggagctgcttaagtcgagtgatcttcttcaatgggacggagcgggtgcggttccctggagagaca  
cttccataaccaggaggagtagCgcgcttcgacagcgacgtgggggagtagcgggcggtagaggagctggggcgg  
cctgtcgccgagtagtggaacagccagaaggacctcctggagcagaagcggggccAggtggacaaTtactgcagac

acaactacggggttggtagagcttcaca (SEQ ID NO: 3 5 3) ;

DRB3\*0210 :

ggggacacccgaccacgtttcttgGagctgcttaagtcigagtgicatttcttcaatgggacggagcgggtgcggt  
tcttggagagaCacttccataaccaggaggagtacGcgcttcgacagcgacgtgggggagtaccgggcggtgac  
5 ggagctggggcgccctgatgccgagtlactggaacagccagaaggacctcttggagcagaagcggggccAggtggac  
aaTtactgcagacacaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 3 5 4) ;

DRB3\*0211 :

ggggacacccgaccacgtttcttggagctgcttaagtcigagtgicatttcttcaatgggacggagcgggtgcggt  
tcttggagagacacttccataaccaggaggagtlacGcgcttcgacagcgacgtgggggagtaccgggcggtgaG  
10 ggagctggggcgccctgatgccgagtlactggaacagccagaaggacAtcttggagcagaagcggggccagggtggac  
aaTtactgcagacacaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 3 5 5) ;

DRB3\*0212 :

cacgtttcttgcagctgcttaagtcigagtgicatttcttcaatgggacggagcgggtgcggttcttggagagaCa  
cttccataaccaggaggagtlacGcgcttcgacagcgacgtgggggagtlaccgggcggtgaGggagctggggcgg  
15 cctgatgccgagtlactggaacagccagaaggacctcttggagcagaagcggggccagggtggacaTtactgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 3 5 6) ;

DRB3\*0213 :

cacgtttcttggagctgcttaagtcigagtgicatttcttcaatgggacggagcgggtgcggttcttggagagaca  
cttccataaccaggaggagtlacGcgcttcgacagcgacgtgggggagtlaccgggcggtgagggagctggggcgg  
20 cctgatgccgagtlactggaacagccagaaggacctcttggagcagaagcggggccagggtggacaattactgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 3 5 7) ;

DRB3\*0214 :

cacgtttcttggagctgcttaagtcigagtgicatttcttcaatgggacggagcgggtgcggttcttggagagaca  
cttccataaccaggaggagtlacGcgcttcgacagcgacgtgggggagtlaccgggcggtgagggagctggggcgg  
25 cctgatgccgagtlactggaacagccagaaggacctcttggagcagaagcggggccagggtggacaattactgcagac  
acaactacggggttgcTtagagcttcacagtcagcggcgag (SEQ ID NO: 3 5 8) ;

DRB3\*0215 :

cacgtttcttggagctgcttaagtcctgagtgctatcttccaatgggacggagcgggtgcggttccctggagagaCa  
cttccataaccaggaggagtagCgcgccttcgacagcgacgtgggggagtagcgggcggtgaGggagctggggcgg  
cctgatgccgagtagtgaacagccagaaggaccttctggagcagaagcggggccAggtggacacctactgcagac  
acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 3 5 9) ;

## 5 DRB3\*0216 :

cacgtttcttggagctgcttaagtcctgagtgctatcttccaatgggacggagcgggtgcggttccctggagagaca  
cttccataaccaggaggagtagcgcgccttcgacagcgacgtgggggagtagcgggcggtgaGggagctggggcgg  
cctgctgcggagCactggaacagccagaaggaccttctggagcagaagcggggccaggtaggacaaTtactgcagac  
acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 3 6 0) ;

## 10 DRB3\*0217 :

cacgtttcttggagctgcttaagtcctgagtgctatcttccaatgggacggagcgggtgcggttccctggagagaca  
cttccataaccaggaggagtagcgcgccttcgacagcgacgtgggggagtagcgggcggtgaGggagctggggcgg  
cctgatgccgagtagtgaacagccagaaggaccttctggagcagaagcggggccaggtaggacaaTtactgcagac  
acaactacggggttggtgagagcttcacagtcagcggcgag (SEQ ID NO: 3 6 1) ;

## 15 DRB3\*030101 :

ggggacacccgaccacgtttcttggagctgcttaagtcctgagtgctatcttccaatgggacggagcgggtgcggt  
tccctggagagatacttccataaccaggaggagtagcgtgcgccttcgacagcgacgtgggggagtagcgggcggtgac  
ggagctggggcggccctgctgccgagtagtgaacagccagaaggaccttctggagcagaagcggggccaggtaggac  
aaTtactgcagacacaactacggggttggtGgagagcttcacagtcagcggcgag (SEQ ID NO: 3 6 2) ;

## 20 DRB3\*030102 :

cacgtttcttggagctgcttaagtcctgagtgctatcttccaatgggacggagcgggtgcggttccctggagagata  
cttccataaccaggaggagtagcgtgcgccttcgacagcgacgtgggggagtagcgggcggtgacggagctggggcgg  
cctgctgccgagtagtgaacagccagaaggaccttctggagcagaagcggggccaggtaggacaaTtactgcagac  
acaactacggCgttggtggagagcttcacagtcagcggcgag (SEQ ID NO: 3 6 3) ;

## 25 DRB3\*0302 :

cacgtttcttggagctgcttaagtcctgagtgctatcttccaatgggacggagcgggtgcggttccctggagagaCa  
cttccataaccaggaggagtagcgtgcgccttcgacagcgacgtgggggagtagcgggcggtgacggagctggggcgg



cctgtcgcgagtcctggaacagccagaaggacctcctggagcagaagcggggccaggctggacaaTtactgcagac  
acaactacggggttgtGg (SEQ ID NO: 3 6 4) ;

DRB3\*0303 :

tttcttggagctgcctaaagtcgagtgctcatttcttcaatgggacggagcgggtgcggttccctggaGagatacttc  
5 cataaccaggaggagtTcgtgcgcttcgacagcgacgtgggggaglacgggcgggtgacggagctggggcggcctg  
tCgccgagtcctggaacagccagaaggacctcctggagcagaagcggggccGggtggacaaTtactgcagacacaa  
ctacggggttggtgagagcttcaca (SEQ ID NO: 3 6 5) ;

DRB4\*010101 :

atgggtgtgtcgaagctccctggaggctccctgtatggcagcgctgacagtgcatttgaCgggtgtgagctccccac  
10 tggctttggctggggacaccaaccacgtttcttggagcaggctaagtgtgagtgctcatttcttcaatgggacgga  
gcgagtggtggaacctgatcagatacatctataaccaagaggagtlacgcgcgtacaacagtgacctgggggagtlac  
caggcgggtgacggagctggggcggcctgacgctgagtlactggaacagccagaaggacctcctggagcggaggcggg  
ccgaggltggacacctactgcagatacaactacggggttgtggagagcttcacagtcagcggcgag (SEQ ID  
NO: 3 6 6) ;

15 DRB4\*0102 :

gagcgagtggtggaacctgatcagatacatctataaccaagaggagtlacgcgcgtacaacagtgacctgggggagtl  
accaggcgggtgacggagctggggcggcctgacgctgagtlactggaacagccagaaggacctcctggagcggaggcg  
ggccgaggltggGcacctactgcagatacaactacggggttgtggagagcttcacagtcagcggcgag (SEQ ID  
NO: 3 6 7) ;

20 DRB4\*010302 :

ggggacaccaaccacgtttcttggagcaggctaagtgtgagtgctcatttcttcaatgggacggagcagtggtgga  
aCctgatcagatacatctataaccaagaggagtlacgcgcgtacaacagtgacctgggggagtlaccaggcggtgac  
ggagctggggcggcctgacgctgagtlactggaacagccagaaggacctcctggagcggaggcgggcccagggtggac  
acctactgcagaTacaactacggggttgtggagagcttcacagtcagcggcgag (SEQ ID NO: 3 6 8) ;

25 DRB4\*010303 :

atgggtgtgtcgaagctccctggaggctccctgtatggcagcgctgacagtgcatttgaCgggtgtgagctccccac  
tggctttggctggggacaccaaccacgtttcttggagcaggctaagtgtgagtgctcatttcttcaatgggacgga

gcgagtggtggaacctgatcagatacatctataaccaagaggagtagcgcgctacaacagtgaacctgggggagtag  
caggcggtagcgagctggggcgccctgacgctgagtagttagaacagccagaaggacctcctggagcggaggcggg  
ccgaggtggacacctatgacagatacaactacggggttgtagagagcttcacagtagcagcggcgag (SEQ ID  
NO: 369) ;

5 DRB4\*010304 :

cacgtttcttggagcaggctaagtgtagtgtagtcttctcaatgggacggagcgagtggtggaacctgatcagata  
catctataaccaagaggagtagcgcgctacaacagttagctgggggagtagcaggcggtagcgagctggggcgg  
cctgacgctgagtagtagaacagccagaaggacctcctggagcggaggcgggcccagggtggacacctactgcagat  
acaactacggggttgtagagagcttcacagtagcagcggcgag (SEQ ID NO: 370) ;

10 DRB4\*0104 :

cacgtttcttggagcaggctaagtgtagtgtagtcttctcaatgggacggagcgagtggtggaacctgatcagata  
catctataaccaagaggagtagcgcgctacaacagttagctgggggagtagcaggcggtagcgagctggggcgg  
cctgacgctgagtagtagaacagccagaaggacctcctggagcggaggcgggcccagggtggacaactactgcagat  
acaactacggggttgtagagagcttcacagtagcagcggcgag (SEQ ID NO: 371) ;

15 DRB4\*0105 :

ttaggagcaggctaagtgtagtgtagtcttctcaatgggacggagcgagtggtggaacctgatcagatacatctata  
accaagaggagtagcgcgctacaacagttagctgggggagtagcaggcggtagcgagctggggcggcctgacgc  
tgagtagtagaacagccagaaggacctcctggagcggaggcgggcccagggtggacacctactgcagacacaactac  
ggggttgtagagag (SEQ ID NO: 372) ;

20 DRB4\*0106 :

cacgtttcttggagcaggctaagtgtagtgtagtcttctcaatgggacggagcgagtggtggaacctgatcagata  
catctataaccaagaggagtagcgcgctacaacagttagctgggggagtagcaggcggtagcgagctggggcgg  
cctgacgctgagtagtagaacagccagaaggacctcctggagcggaggcgggcccagggtggacacctactgcagat  
acaactacggggttgtagagagcttcacagtagcagcggcgag (SEQ ID NO: 373) ;

25 DRB4\*0201N :

ggtagtagctccccactggctttggctggggacacccAaccacgtttcttggagcaggctaagtgtagtgtagt  
tctctcaatgggacggagcctgatcagatacatctataaccaagaggagtagcgcgctacaacagttagctggg

gagtagcaggcggtgacggagctggggcgccctgacgctgagtagtggaacagccagaaggacctccaggagcgga  
ggcgggcccagggtggacacctactgcagatacaactacggggctgtGgagagcttcacagtcagcggcgag (SEQ  
ID NO: 374) ;

DRB5\*010101 :

5 atgggtgtgtcgaagctccctggaggctccctacatggcaaAgctgacagtgacactgaatgggtgtgagctccccac  
tggctttggctggggacacccgaccagctttcttcgagcaggataagtagtagtgcatttttcaacgggacggga  
gcgggtgcggttcttcacagagacatctataaccaagaggaggacttgcgcttcgacagcgacgtgggggagtag  
cgggcggtgacggagctggggcgccctgacgctgagtagtggaacagccagaaggacttccaggaaacagggcgcg  
ccgcggtggacacctactgcagacacaactacggggctggtagagcttcacagtcagcggcgag (SEQ ID

10 NO: 375) ;

DRB5\*010102 :

cacgtttcttcgagcaggataagtagtagtgcatttttcaacgggacggagcgggtgcggttcttcacagaga  
catctataaccaagaggaggacttgcgcttcgacagcgacgtgggggagtagcggcggtgacggagctggggcg  
cctgacgctgagtagtggaacagccagaaggacttccaggaaGacaggcgccggtggacacctactgcagac  
15 acaactacggggctggtagagcttcaca (SEQ ID NO: 376) ;

DRB5\*0102 :

ggggacacccgaccagctttcttcgagcaggataagtagtagtgcatttttcaacgggacggagcgggtgcggt  
tcttcacagaggcatctataaccaagaggagAACgtgcgcttcgacagcgacgtgggggagtagcggcggtgac  
ggagctggggcgccctgacgctgagtagtggaacagccagaaggacttccaggaaGacaggcgCgccggtggac

20 acctactgcagacacaactacggggctggtagagcttcacagtcagcggcgag (SEQ ID NO: 377) ;

DRB5\*0103 ;

ttgcagcaggataagtagtagtgcatttttcaacgggacggagcgggtgcggttcttcacagaGgcatctata  
accaagaggagaacgtgcgcttcgacagcgacgtgggggagtagcggcggtgacggagctggggcgccctgacgc  
tagtagtagtggaacagccagaaggacttccaggaaGacaggcgCgccggtggacacctactgcagacacaactac

25 ggggttggtagagcttcacag (SEQ ID NO: 378) ;

DRB5\*0104 :

ggggacacccgaccagctttcttcgagcaggataagtagtagtgcatttttcaacgggacggagcgggtgcggt

tccatgcacagagacatctataaccaagaggaggacTtgcgcttcgacagcgacgtgggggagtagcgggaggtagac  
ggagctggggcgccctgacgctgagtagtggaaacagccagaaggacttccatggaagacaggcggggccTggtggac  
acctactgcagacacaactacggggttggtagagagcttcacagtcagcggcgag (SEQ ID NO: 379) ;  
DRB5\*0105 :

- 5 ccacgtttcttgcagcaggataagtagtagtgcatttcttcaacgggacggagcgggtgcggttccatgcacagag  
acatctataaccaagaggagGacgtgcgcttcgacagcgacgtgggggagtagcgggaggtagcggagctggggcg  
gctgacgctgagtagtggaaacagccagaaggacTtccatggaGacaggcgCgccgaggtagacacctactgcaga  
cacaactacggggttggtagagagcttcacagtcagcgg (SEQ ID NO: 380) ;

DRB5\*0106 :

- 10 cacgtttcttgcagcaggataagtagtagtgcatttcttcaacgggacggagcgggtgcggttccatgcacagaga  
catctataaccaagaggaggacTtgcgcttcgacagcgacgtgggggagtagcgggaggtagcggagctggggcgg  
cctgacgctgagtagtggaaacagccagaaggacatccatggagcaggcggggccgaggtagacacctactgcagac  
acaactacggggttggtagagagcttcacagtcagcggcga (SEQ ID NO: 381) ;

DRB5\*0107 :

- 15 cacgtttcttgcagcaggataagtagtagtgcatttcttcaacgggacggagcgggtgcggttccatgcacagaga  
catctataaccaagaggaggacTtgcgcttcgacagcgacgtgggggagtagcgggaggtagcggagctggggcgg  
cctgacgctgagtagtggaaacagccagaaggacAtccatggaGacaggcgCgccgaggtagacacctactgcagac  
acaactacggggttggtag (SEQ ID NO: 382) ;

DRB5\*0109 :

- 20 cacgtttcttgcagcaggataagtagtagtgcatttcttcaacgggacggagcgggtgcggttccatgcacagaga  
catctataaccaagaggaggacttgcgcttcgacagcgacgtgggggagtagcgggaggtagcggagctggggcgg  
cctgacgctgagtagtggaaacagccagaaggacttccatggaAacaggcgCgccgaggtagacacctactgcagac  
acaactacggggttggtag (SEQ ID NO: 383) ;

DRB5\*0110N :

- 25 cacgtttcttgcagcaggataagtagtagtgcatttcttcaacgggacggagcgggtgcggttccatgcacagaGg  
catctataaccaagaggagAacgtgcgcttcgacagcgacgtgggggagtagcgggaggtagcggagctggggcgg  
cctgacgctgagtagtggaaacagccagaaggacTtccatggaGacaggcgCgccgaggtagacacctactgca...c

acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 384) ;

DRB5\*0111 :

cacgtttcttgcagcaggataagtaagtagtgatcttcttcaacgggacggagcgggtgcggttccctgcacagaga  
catctataaccaagaggaggacTtgcgcttcgacagcgacgtgggggagtagcggggtgacggagctggggcgg

5 cctgacgctgagtagtggaaacagccagaaggacatcctggagcaggCgcgggccggtggacacctactgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 385) ;

DRB5\*0112 :

cacgtttcttgcagcaggataagtaagtagtgatcttcttcaacgggacggagcgggtgcggttccctgcacagaga  
catctataaccaagaggaggacTtgcgcttcgacagcgacgtgggggagtagcggggtgacggagctggggcgg

10 cctgacgcccagtagtCctggaaacagccagaaggacttctggagcGgaggcgggccgaggtggacaccgtGtgcagac  
acaactacggggttggtagagcttcacagtcagcggcgag (SEQ ID NO: 386) ;

DRB5\*0202 :

atgggtgtgtcgaagctccctggaggttcttAcatggcagtagctgacagtgacactgaatgggtgtgagctccccac  
tggctttggctggggacacccgacctgtttcttgcagcaggataagtaagtagtgatcttcttcaacgggacgga

15 gcgggtgcggttccctgcacagaggcatctataaccaagaggagaacgtgcgcttcgacagcgacgtgggggagtag  
cggggtgacggagctggggcggctgacgctgagtagtggaaacagccagaaggacatcctggagcaggcgcggg  
ccgcggtggacacctactgcagacacaactacggggctgtGgagagcttcacagtcagcggcgag (SEQ ID  
NO: 387) ;

DRB5\*0203 :

20 ttcttgcagcaggataagtaagtagtgatcttcttcaacgggacggagcgggtgcggttccctgcacagaGgcatt  
tataaccaagaggagAacgtgcgcttcgacagcgacgtgggggagtagcggggtgacggagctggggcggctg  
acgtgagtagtggaaacagccagaaggacAtcctggagcagCgcgggccggtggacacctactgcagacacaa  
ctacggggttggtagagcttcacagtcagcgg (SEQ ID NO: 388) ;

DRB5\*0204 :

25 catgtttcttgcagcaggataagtaagtagtgatcttcttcaacgggacggagcgggtgcggttccctgcacagaGg  
catctataaccaagaggagaacgtgcgcttcgacagcgacgtgggggagtagcggggtgacggagctggggcgg  
cctgacgctgagtagtggaaacagccagaaggacTtctggagcaggCgcgggccggtggacacctactgcagac

acaactacggggctgtGgagagcttcaca (SEQ ID NO: 3 8 9) ;

DRB5\*0205 :

catgtttcttgcagcaggataagtaagtgatgtcatgtttcttcaacgggacggagcgggtgcggttccgtgcacagaGg

catctataaccaagaggagAacgtgcgcttcgacagcgacgtgggggagtagcgggcgggtgacggagctggggcgg

5 cctgacgctgagtactggaacagccagaaggacctcctggagcagaggcgggccggtggacacctactgcagac

acaactacggggctgtGgagagcttcacagtgcagcggcgag (SEQ ID NO: 3 9 0)

In the following, Probe List DR1 and 2 are  
shown in Tables 21-1 to 21-8 and Tables 22-1 to 22-7  
10 respectively. Allele-Probe Lists 1 and 2 are shown  
in Tables 23-1 to 23-13 and Tables 24-1 to 24-13  
respectively.

Table 21-1

Probe No.	Base Sequence
0	g gtg cgg ttg ctg gaA ( SEQ ID No: 391)
1	g cgg ttg ctg gaa aga T ( SEQ ID No: 392)
2	c tat aac caa gag gag tC ( SEQ ID No: 393)
3	ctg ggg cgg cct gaT ( SEQ ID No: 394)
4	ggg cgg cct gat gcC ( SEQ ID No: 395)
5	cac aac tac ggg gtt gG ( SEQ ID No: 396)
6	c atc tat aac caa gag gaA ( SEQ ID No: 397)
7	c gcg gtg gac acc taT ( SEQ ID No: 398)
8	ga cac aac tac ggg gC ( SEQ ID No: 399)
9	ag agg cgg gcc gcC ( SEQ ID No: 400)
10	g aac agc cag aag gac A ( SEQ ID No: 401)
11	g gac atc ctg gaa gac G ( SEQ ID No: 402)
12	gac atc ctg gaa gac gA ( SEQ ID No: 403)
13	g gcc gcg gtg gac aaT ( SEQ ID No: 404)
14	ac aac tac ggg gtt gtG ( SEQ ID No: 405)
15	c ttc gac agc gac gtg A ( SEQ ID No: 406)
16	c ctc ctg gag cag gC ( SEQ ID No: 407)
17	ca cgt ttc ttg tgg G ( SEQ ID No: 408)
18	tc tat aac caa gag gag tA ( SEQ ID No: 409)
19	gac ctc ctg gag cag G ( SEQ ID No: 410)
20	gac ctc ctg gag cag aA ( SEQ ID No: 411)
21	g gag cgg gtg cgg tA ( SEQ ID No: 412)
22	c ctg gac aga tac ttc C ( SEQ ID No: 413)
23	c cat aac cag gag gag A ( SEQ ID No: 414)
24	c cat aac cag gag gag aA ( SEQ ID No: 415)
25	gc gac gtg ggg gag tT ( SEQ ID No: 416)
26	g cag aag cgg ggc cG ( SEQ ID No: 417)
27	g ggc cgg gtg gac aA ( SEQ ID No: 418)
28	g ggc cgg gtg gac aaT ( SEQ ID No: 419)
29	ca cgt ttc ttg gA ( SEQ ID No: 420)
30	g gtg cgg ttc ctg gaG ( SEQ ID No: 421)

Table 21-2

Probe No.	Base Sequence
31	c ctg gag aga tac ttc C ( SEQ ID No: 4 2 2)
32	c aga tac ttc cat aac caG ( SEQ ID No: 4 2 3)
33	tt ggt gag agc ttc acG ( SEQ ID No: 4 2 4)
34	g gtg cgg tac ctg gaC ( SEQ ID No: 4 2 5)
35	g ggg cgg cct gat gA ( SEQ ID No: 4 2 6)
36	ggg cgg cct gat gaG ( SEQ ID No: 4 2 7)
37	c aga tac ttc cat aac cG ( SEQ ID No: 4 2 8)
38	ctg ggg cgg cct gC ( SEQ ID No: 4 2 9)
39	ag cag aag cgg ggc C ( SEQ ID No: 4 3 0)
40	g cag aag cgg ggc cA ( SEQ ID No: 4 3 1)
41	gg ggc cag gtg gac aA ( SEQ ID No: 4 3 2)
42	ctg ggg cgg cct agC ( SEQ ID No: 4 3 3)
43	gg cct gat gcc gag tC ( SEQ ID No: 4 3 4)
44	gac gtg ggg gag ttc T ( SEQ ID No: 4 3 5)
45	gt ttc ttg gag tac tct aC ( SEQ ID No: 4 3 6)
46	g gtg cgg ttc ctg gaC ( SEQ ID No: 4 3 7)
47	g tac cgg gcg gtg aG ( SEQ ID No: 4 3 8)
48	g ggc cag gtg gac aaT ( SEQ ID No: 4 3 9)
49	ttc gac agc gac gtg C ( SEQ ID No: 4 4 0)
50	c cat aac cag gag gag tT ( SEQ ID No: 4 4 1)
51	c ctg gac aga tac ttc G ( SEQ ID No: 4 4 2)
52	c cat aac cag gag gag tA ( SEQ ID No: 4 4 3)
53	atg gtg tgt ctg aag T ( SEQ ID No: 4 4 4)
54	ga tac ttc tat cac caa gaA ( SEQ ID No: 4 4 5)
55	tc ttg gag cag gtt aaa C ( SEQ ID No: 4 4 6)
56	c tat cac caa gag gag tA ( SEQ ID No: 4 4 7)
57	g cag agg cgg gcc gA ( SEQ ID No: 4 4 8)
58	ggg cgg cct gac gcT ( SEQ ID No: 4 4 9)
59	c ttg gag cag gtt aaa cA ( SEQ ID No: 4 5 0)
60	ctg gac aga tac ttc tat C ( SEQ ID No: 4 5 1)



Table 21-3

Probe No.	Base Sequence
61	g ctg ggg cgg cct aG ( SEQ ID No: 452)
62	a gag gag tac gtg cgG ( SEQ ID No: 453)
63	gc ttc aca gtg cag cgA ( SEQ ID No: 454)
64	c ctc ctg gag cag agA ( SEQ ID No: 455)
65	t ttc ttg gag cag gtt aaA ( SEQ ID No: 456)
66	a gac agg cgg gcc cT ( SEQ ID No: 457)
67	g aac agc cag aag gac T ( SEQ ID No: 458)
68	ag gac ttc ctg gaa gaC ( SEQ ID No: 459)
69	gg cgg cct gat gcc C ( SEQ ID No: 460)
70	c ggg gtt gtg gag agA ( SEQ ID No: 461)
71	g gac ctc ctg gag cG ( SEQ ID No: 462)
72	ctg ggg cgg cct gat A ( SEQ ID No: 463)
73	ag tac cgg gcg gtg aT ( SEQ ID No: 464)
74	g ggg gag tac cgg gT ( SEQ ID No: 465)
75	g cag agg cgg gcc C ( SEQ ID No: 466)
76	g cag agg cgg gcc cT ( SEQ ID No: 467)
77	tc ctg gag cag agg cA ( SEQ ID No: 468)
78	caa gag gag tac gtg cA ( SEQ ID No: 469)
79	c ttg gag cag gtt aaa cC ( SEQ ID No: 470)
80	gac ctc ctg gaa gac G ( SEQ ID No: 471)
81	gac ctc ctg gaa gac gA ( SEQ ID No: 472)
82	gac atc ctg gag cag aA ( SEQ ID No: 473)
83	agc gac gtg ggg gaC ( SEQ ID No: 474)
84	g ggg cgg cct gat gG ( SEQ ID No: 475)
85	tc tat cac caa gag gag A ( SEQ ID No: 476)
86	c tat cac caa gag gag aA ( SEQ ID No: 477)
87	g gct ggg gac acc cA ( SEQ ID No: 478)
88	g gac agg cgg ggc C ( SEQ ID No: 479)
89	c cag gtg gac acc gtG ( SEQ ID No: 480)
90	tc ctg tgg cag ggt aaA ( SEQ ID No: 481)

Table 21-4

Probe No.	Base Sequence
91	g gcg gtg acg gag ctA ( SEQ ID No: 482)
92	g cct gtc gcc gag tC ( SEQ ID No: 483)
93	gtg cag ttc ctg gaa agT ( SEQ ID No: 484)
94	ag tcc tgg aac agc cG ( SEQ ID No: 485)
95	gg cgg cct gct gcG ( SEQ ID No: 486)
96	gtg acg gag cta ggg T ( SEQ ID No: 487)
97	c tct acg ggt gag tgt T ( SEQ ID No: 488)
98	cgg ttc ctg gac aga taT ( SEQ ID No: 489)
99	gc tcc tgc atg gca gT ( SEQ ID No: 490)
100	g tac cgg gcg gtg acA ( SEQ ID No: 491)
101	cac aac tac ggg gtt gT ( SEQ ID No: 492)
102	gtt gtt gag agc ttc acG ( SEQ ID No: 493)
103	tt gtg gag agc ttc acG ( SEQ ID No: 494)
104	g ctg ggg cgg cct gT ( SEQ ID No: 495)
105	gg cct gct gcg gag C ( SEQ ID No: 496)
106	gt ttc ttg gag tac tct aG ( SEQ ID No: 497)
107	gg cct gat gcg gag C ( SEQ ID No: 498)
108	tc tat aac caa gag gag G ( SEQ ID No: 499)
109	ag gac atc ctg gaa gaC ( SEQ ID No: 500)
110	g ctg ggg cgg cct aT ( SEQ ID No: 501)
111	c ttg gag tac tct acg tC ( SEQ ID No: 502)
112	gt ttc ttg gag tac tct aT ( SEQ ID No: 503)
113	c aac tac ggg gct gtG ( SEQ ID No: 504)
114	ct gtg gag agc ttc acG ( SEQ ID No: 505)
115	g agc ttc aca gtg cag A ( SEQ ID No: 506)
116	ctg gag cgg agg cgT ( SEQ ID No: 507)
117	g ttg ctg gaa aga cgc G ( SEQ ID No: 508)
118	ctg gag cgg agg cgC ( SEQ ID No: 509)
119	g aag gac ttc ctg gaa G ( SEQ ID No: 510)
120	c ctg gaa gac agg cgC ( SEQ ID No: 511)

Table 21-5

Probe No.	Base Sequence
121	t gag tgt cat ttc ttc aaC ( SEQ ID No: 512)
122	gac ttc ctg gaa gac gA ( SEQ ID No: 513)
123	c ttg gag tac tct acg G ( SEQ ID No: 514)
124	g gac ctc ctg gaa gaC ( SEQ ID No: 515)
125	g gac ttc ctg gaa gac G ( SEQ ID No: 516)
126	tc tat aac caa gag gag tT ( SEQ ID No: 517)
127	c aga tac ttc tat aac caG ( SEQ ID No: 518)
128	c tat aac cag gag gag tT ( SEQ ID No: 519)
129	at aac caa gag gag gac T ( SEQ ID No: 520)
130	cgg agg cgg gcc gA ( SEQ ID No: 521)
131	cc gag gtg gac acc taT ( SEQ ID No: 522)
132	aa gac agg cgg gcc C ( SEQ ID No: 523)
133	ttg gag tac tct acg tC ( SEQ ID No: 524)
134	gag tac tct acg tct gaG ( SEQ ID No: 525)
135	cag aag gac ttc ctg gaA ( SEQ ID No: 526)
136	g gcc gcg gtg gac aA ( SEQ ID No: 527)
137	ttc tat aac caa gag gag A ( SEQ ID No: 528)
138	tc tat aac caa gag gag aA ( SEQ ID No: 529)
139	ca cgt ttc ttg gag cT ( SEQ ID No: 530)
140	cgg cct gat gag gag C ( SEQ ID No: 531)
141	a gac agg cgg gcc gT ( SEQ ID No: 532)
142	g cgg cct gat gag gaC ( SEQ ID No: 533)
143	g cgg cct gat gag gG ( SEQ ID No: 534)
144	g ttc cgg gcg gtg aG ( SEQ ID No: 535)
145	gc tcc tgc atg gca gtT ( SEQ ID No: 536)
146	ttg gct ggg gac acc A ( SEQ ID No: 537)
147	g gag cgg gtg cgg ttA ( SEQ ID No: 538)
148	c cat aac cag gag gag C ( SEQ ID No: 539)
149	cag aag gac atc ctg gG ( SEQ ID No: 540)
150	gag cgg gtg cgg ttC ( SEQ ID No: 541)

Table 21-6

Probe No.	Base Sequence
151	g gaa gac gag cgg gcT ( SEQ ID No: 542)
152	c ctg gaa gac gag cgC ( SEQ ID No: 543)
153	g gac atc ctg gaa gac aA ( SEQ ID No: 544)
154	a cgt ttc ttg gag tac tC ( SEQ ID No: 545)
155	gg ttc ctg gac aga tac T ( SEQ ID No: 546)
156	ac atc ctg gag cag gC ( SEQ ID No: 547)
157	cac aac tac ggg gtt gA ( SEQ ID No: 548)
158	g aga tac ttc cat aac caG ( SEQ ID No: 549)
159	c tgc aga cac aac tac C ( SEQ ID No: 550)
160	t aac cag gag gag aac C ( SEQ ID No: 551)
161	ac gtg ggg gag ttc cT ( SEQ ID No: 552)
162	ctg ggg cgg cct gtC ( SEQ ID No: 553)
163	gg gag ttc cgg gcg T ( SEQ ID No: 554)
164	ca cgt ttc ttg gag tac T ( SEQ ID No: 555)
165	tct acg tct gag tgt caA ( SEQ ID No: 556)
166	ggg cgg cct gat gcT ( SEQ ID No: 557)
167	t ttc ttg gag tac tct aC ( SEQ ID No: 558)
168	gac atc ctg gag cag G ( SEQ ID No: 559)
169	g acg gag cgg gtg cA ( SEQ ID No: 560)
170	g gcc gag gtg gac aaT ( SEQ ID No: 561)
171	ttg gag tac cct acg tC ( SEQ ID No: 562)
172	t aac cag gag gag ttc C ( SEQ ID No: 563)
173	gg gcc gag gtg gac G ( SEQ ID No: 564)
174	c tcc cca ctg gct ttg T ( SEQ ID No: 565)
175	gc aga cac aac tac ggA ( SEQ ID No: 566)
176	cac aac tac gga gtt gtG ( SEQ ID No: 567)
177	g tgg cag cct aag agG ( SEQ ID No: 568)
178	tg gac aga tac ttc tat aaT ( SEQ ID No: 569)
179	cgg ttc ctg gac aga C ( SEQ ID No: 570)
180	ac ttc ctg gag cag gC ( SEQ ID No: 571)

Table 21-7

Probe No.	Base Sequence
181	g gag ttc cgg gcg gC ( SEQ ID No: 572)
182	c tgg aac agc cag aag A ( SEQ ID No: 573)
183	ac gtg ggg gag ttc cA ( SEQ ID No: 574)
184	c tgg aac agc ca ggg gac A ( SEQ ID No: 575)
185	tc ctg gaa gac agg gC ( SEQ ID No: 576)
186	g cgg gtg cgg ttc cC ( SEQ ID No: 577)
187	c tat aac cag gag gag aA ( SEQ ID No: 578)
188	cgt ttc ttg gag ctg cG ( SEQ ID No: 579)
189	c tcc cga ctg gct ttC ( SEQ ID No: 580)
190	ca cgt ttc ttg gag ctg T ( SEQ ID No: 581)
191	cgt ttc ttg gag ctg tG ( SEQ ID No: 582)
192	g gtg cgg tac ctg gaG ( SEQ ID No: 583)
193	gt ttc tgc gag ctg cG ( SEQ ID No: 584)
194	cgg gtg cgg tac ctg A ( SEQ ID No: 585)
195	ac cag gag gag tac gC ( SEQ ID No: 586)
196	c cag gag gag ttc ctg A ( SEQ ID No: 587)
197	ca cgt ttc ttg G ( SEQ ID No: 588)
198	cgg ttc ctg gag aga C ( SEQ ID No: 589)
199	gtg gac aat tac tgc agG ( SEQ ID No: 590)
200	ggg cgg cct gat gcG ( SEQ ID No: 591)
201	aga cac ttc cat aac caG ( SEQ ID No: 592)
202	ac cag gag gag aac gC ( SEQ ID No: 593)
203	g gag cgg gtg cgg C ( SEQ ID No: 594)
204	cac aac tac ggg gti gC ( SEQ ID No: 595)
205	gc aga cac aac tac ggC ( SEQ ID No: 596)
206	g ctg aca gtg aca ttg aC ( SEQ ID No: 597)
207	cgg gcc gag gtg gG ( SEQ ID No: 598)
208	ag tgt gag tgt cat ttc C ( SEQ ID No: 599)
209	g gag cga gtg tgg aaC ( SEQ ID No: 600)
210	g gac acc tac tgc aga T ( SEQ ID No: 601)

Table 21-8

Probe No.	Base Sequence
211	cg cgc tac aac agt gaT ( SEQ ID No: 602)
212	gg gcc gag gtg gac aA ( SEQ ID No: 603)
213	tg gac aac tac tgc aga T ( SEQ ID No: 604)
214	acg gag cga gtg tgg A ( SEQ ID No: 605)
215	a ggt tcc tac atg gca aA ( SEQ ID No: 606)
216	ca cgt ttc ttg C ( SEQ ID No: 607)
217	atc tat aac caa gag gag A ( SEQ ID No: 608)
218	cgg ttc ctg cac aga G ( SEQ ID No: 609)
219	gac ttc ctg gaa gac aC ( SEQ ID No: 610)
220	c ctg gaa gac acg cgC ( SEQ ID No: 611)
221	g aag gac atc ctg gaa G ( SEQ ID No: 612)
222	ag aag gac ttc ctg gaa A ( SEQ ID No: 613)
223	g cct gac gcc gag tC ( SEQ ID No: 614)
224	ag gac ttc ctg gag cG ( SEQ ID No: 615)
225	c gag gtg gac acc gtG ( SEQ ID No: 616)
226	ctc cct gga ggt tcc tA ( SEQ ID No: 617)

Table 22-1

Probe No.	Base Sequence
0	g ttg ctg gaA aga tgc at ( SEQ ID No: 618)
1	ctg gaa aga Tgc atc tal a ( SEQ ID No: 619)
2	gag gag tCc gtg cgc ( SEQ ID No: 620)
3	cgg cct gaT gcc gag ( SEQ ID No: 621)
4	cct gat gcC gag tac tg ( SEQ ID No: 622)
5	c ggg gtt gGt gag agc ( SEQ ID No: 623)
6	caa gag gaA tcc gtg cg ( SEQ ID No: 624)
7	g gac acc taT tgc aga ca ( SEQ ID No: 625)
8	c tac ggg gCt gtg gag ( SEQ ID No: 626)
9	gg gcc gcC gtg gac ( SEQ ID No: 627)
10	cag aag gac Atc ctg gaa ( SEQ ID No: 628)
11	g gaa gac Gag cgg gc ( SEQ ID No: 629)
12	gaa gac gAg cgg gcc ( SEQ ID No: 630)
13	g gtg gac aaT tac tgc ag ( SEQ ID No: 631)
14	ggg gtt gtG gag agc t ( SEQ ID No: 632)
15	c gac gtg Agg gag tac ( SEQ ID No: 633)
16	gag cag gCg cgg gc ( SEQ ID No: 634)
17	ttc ttg tgg Gag ctt aag ( SEQ ID No: 635)
18	a gag gag tAc gtg cgc ( SEQ ID No: 636)
19	gag cag Gcg cgg gc ( SEQ ID No: 637)
20	gag cag aAg cgg gcc ( SEQ ID No: 638)
21	xc acc Aga c ( SEQ ID No: 639)
22	g gtg cgg tAc ctg gac ( SEQ ID No: 640)
23	g gtg gac aAc tac tgc a ( SEQ ID No: 641)
24	cgg ggc cGg gtg ga ( SEQ ID No: 642)
25	g ttc ctg gaG aga tac tt ( SEQ ID No: 643)
26	aga tac ttc Cat aac cag g ( SEQ ID No: 644)
27	g gag gag Aac gtg cgc ( SEQ ID No: 645)
28	g gag gag aAc gtg cgc ( SEQ ID No: 646)
29	cat aac caG gag gag tc ( SEQ ID No: 647)
30	ggg gag tTc cgg gcg ( SEQ ID No: 648)

Table 22-2

Probe No.	Base Sequence
31	agc ttc acG gtg cag c ( SEQ ID No: 649)
32	g tac ctg gaC aga tac tt ( SEQ ID No: 650)
33	g cct gat gAg gag tac t ( SEQ ID No: 651)
34	cct gat gaG gag tac tg ( SEQ ID No: 652)
35	c cat aac cGg gag gag ( SEQ ID No: 653)
36	cgg cct gCt gcg gag ( SEQ ID No: 654)
37	g cgg ggc Cag gtg ga ( SEQ ID No: 655)
38	cgg ggc cAg gtg gac ( SEQ ID No: 656)
39	cgg cct aGc gcc gag ( SEQ ID No: 657)
40	cgg cct agC gcc gag ( SEQ ID No: 658)
41	t gcc gag tCc tgg aac ( SEQ ID No: 659)
42	g gag ttc Tgg gcg gtg ( SEQ ID No: 660)
43	ag tac tct aCg tct gag t ( SEQ ID No: 661)
44	g ttc ctg gaC aga tac tt ( SEQ ID No: 662)
45	gcg gtg aGg gag ctg ( SEQ ID No: 663)
46	c gac gtg Cgg gag ttc ( SEQ ID No: 664)
47	ag aag gac ATc ctg gag ( SEQ ID No: 665)
48	g gag gag tTc gtg cgc ( SEQ ID No: 666)
49	aga tac ttc Gat aac cag g ( SEQ ID No: 667)
50	c cat aac caG gag gag ta ( SEQ ID No: 668)
51	g gag gag tAc gtg cgc ( SEQ ID No: 669)
52	gt ctg aag Ttc cct gga ( SEQ ID No: 670)
53	t cac caa gaA gag tac gt ( SEQ ID No: 671)
54	cag gtt aaa Cat gag tgt c ( SEQ ID No: 672)
55	cgg gcc gAg gtg gac ( SEQ ID No: 673)
56	cct gac gCt gag tac tg ( SEQ ID No: 674)
57	ag gtt aaa cAt gag tgt ca ( SEQ ID No: 675)
58	tac ttc tat Cac caa gag g ( SEQ ID No: 676)
59	tac gtg cgG ttc gac ag ( SEQ ID No: 677)
60	gag cag agA cgg gcc ( SEQ ID No: 678)



Table 22-3

Probe No.	Base Sequence
61	g cag gtt aaA cat gag tg ( SEQ ID No: 679)
62	cgg gcc cTg gtg gac ( SEQ ID No: 680)
63	cag aag gac Ttc ctg gaa ( SEQ ID No: 681)
64	ctg gaa gaC agg cgg g ( SEQ ID No: 682)
65	ct gat gcc Cag tac tgg ( SEQ ID No: 683)
66	t gtg gag agA ttc aca gt ( SEQ ID No: 684)
67	ctg gag cGg agg cgg ( SEQ ID No: 685)
68	g cgg gcc Ctg gtg ga ( SEQ ID No: 686)
69	gg cct gat Acc gag tac ( SEQ ID No: 687)
70	g gcg gtg aTg gag ctg ( SEQ ID No: 688)
71	g tac cgg gTg gtg acg ( SEQ ID No: 689)
72	cag agg cAg gcc gcg ( SEQ ID No: 690)
73	g tac gtg cAc ttc gac a ( SEQ ID No: 691)
74	cag gtt aaa Cct gag tgt ( SEQ ID No: 692)
75	ag gtt aaa cCt gag tgt c ( SEQ ID No: 693)
76	gtg ggg gaC tac cgg ( SEQ ID No: 694)
77	g cct gat gGc gag tac ( SEQ ID No: 695)
78	a gag gag Aac gtg cgc ( SEQ ID No: 696)
79	a gag gag aAc gtg cgc ( SEQ ID No: 697)
80	xacc cAa c ( SEQ ID No: 698)
81	gac acc gTG tgc aga c ( SEQ ID No: 699)
82	g cag ggt aaA tat aag tgt ( SEQ ID No: 700)
83	acg gag ctA ggg cgg ( SEQ ID No: 701)
84	c gcc gag tCc tgg aac ( SEQ ID No: 702)
85	c ctg gaa agT ctc ttc ta ( SEQ ID No: 703)
86	g aac agc cGg aag gac ( SEQ ID No: 704)
87	cct gct gcG gag tac t ( SEQ ID No: 705)
88	g cta ggg Tgg cct gtc ( SEQ ID No: 706)
89	ggt gag tgt Tat ttc ttc a ( SEQ ID No: 707)
90	tg gac aga taT ttc tat aac ( SEQ ID No: 708)

Table 22-4

Probe No.	Base Sequence
91	g tgt ctg aGg ctc cct ( SEQ ID No: 709)
92	gcg gtg acA gag ctg g ( SEQ ID No: 710)
93	c ggg gtt gTt gag agc ( SEQ ID No: 711)
94	cgg cct gTt gcc gag ( SEQ ID No: 712)
95	t gcg gag Cac tgg aac ( SEQ ID No: 713)
96	g tac tct aCg ggt gag t ( SEQ ID No: 714)
97	cgg cct gCt gcc gag ( SEQ ID No: 715)
98	g tac tct aGg ggt gag t ( SEQ ID No: 716)
99	a gag gag Gac gtg cgc ( SEQ ID No: 717)
100	cgg cct aTc gcc gag ( SEQ ID No: 718)
101	c tct acg tCt gag tgt c ( SEQ ID No: 719)
102	ag tac tct aTg ggt gag t ( SEQ ID No: 720)
103	ggg gct gtG gag agc ( SEQ ID No: 721)
104	gtg cgg taT ctg cac ag ( SEQ ID No: 722)
105	gg agg cgT gcc gcg ( SEQ ID No: 723)
106	gaa aga cgc Gtc cat aac ( SEQ ID No: 724)
107	gg agg cgC gcc gcg ( SEQ ID No: 725)
108	c ctg gaa Gac agg cgc ( SEQ ID No: 726)
109	ctg gaa gaC agg cgc g ( SEQ ID No: 727)
110	ac agg cgC gcc gcg ( SEQ ID No: 728)
111	ttc ttc aaC ggg acg ga ( SEQ ID No: 729)
112	ac tct acg Ggt gag tgt ( SEQ ID No: 730)
113	c cat aac caG gag gag aa ( SEQ ID No: 731)
114	c cat aac caG gag gag tt ( SEQ ID No: 732)
115	a gag gag tTc gtg cgc ( SEQ ID No: 733)
116	c tat aac caG gag gag tt ( SEQ ID No: 734)
117	g gag gac Ttg cgc ttc ( SEQ ID No: 735)
118	c ctg gaa Gac agg cgg ( SEQ ID No: 736)
119	t acg tct gaG tgt cat ttc ( SEQ ID No: 737)
120	ttc ctg gaA gac agg cg ( SEQ ID No: 738)

Table 22-5

Probe No.	Base Sequence
121	tc ttg gag cTg ctt aag t ( SEQ ID No: 739)
122	g cct gat gAg gag cac ( SEQ ID No: 740)
123	at gag gag Cac tgg aac ( SEQ ID No: 741)
124	cgg gcc gTg gtg gac ( SEQ ID No: 742)
125	t gat gag gaC tac tgg aa ( SEQ ID No: 743)
126	t gat gag gGg tac tgg a ( SEQ ID No: 744)
127	c atg gca gtT ctg aca gt ( SEQ ID No: 745)
128	gtg cgg ttA ctg gag ag ( SEQ ID No: 746)
129	g gag gag Ctc ctg cg ( SEQ ID No: 747)
130	c atc ctg gGa gac agg ( SEQ ID No: 748)
131	gtg cgg ttC ctg gag a ( SEQ ID No: 749)
132	gag cgg gcT gcg gtg ( SEQ ID No: 750)
133	gaa gac gAg cgc gcc ( SEQ ID No: 751)
134	ac gag cgC gcc gcg ( SEQ ID No: 752)
135	ctg gaa gaC aag cgg g ( SEQ ID No: 753)
136	g gaa gac aAg cgg gcc ( SEQ ID No: 754)
137	g gag tac tCt acg tct g ( SEQ ID No: 755)
138	gac aga tac Ttc tat aac c ( SEQ ID No: 756)
139	c ggg gtt gAt gag agc ( SEQ ID No: 757)
140	ac aac tac Cgg gtt gtg ( SEQ ID No: 758)
141	cgg cct gTc gcc gag ( SEQ ID No: 759)
142	g gag aac Ctg cgc ttc ( SEQ ID No: 760)
143	g gag ttc cTg gcg gtg ( SEQ ID No: 761)
144	cgg cct gtC gcc gag ( SEQ ID No: 762)
145	c cgg gcg Ttg acg ga ( SEQ ID No: 763)
146	ttg gag tac Tct acg tct ( SEQ ID No: 764)
147	ct gag tgt caA ttc ttc aat ( SEQ ID No: 765)
148	cct gat gcT gag tac tg ( SEQ ID No: 766)
149	gt ttc ttg gAg tac tct ac ( SEQ ID No: 767)
150	g cgg gtg cAg ttc ctg ( SEQ ID No: 768)

Table 22-6

Probe No.	Base Sequence
151	c gac gtg Cgg gag tac ( SEQ ID No: 769)
152	c cct acg tCt gag tgt c ( SEQ ID No: 770)
153	g gag gag tTc ctg cgc ( SEQ ID No: 771)
154	g gag ttc CtG cgc ttc ( SEQ ID No: 772)
155	g gtg gac Gcc tat tgc ( SEQ ID No: 773)
156	g gct ttg Tct ggg gac ( SEQ ID No: 774)
157	c aac tac ggA gtt gtg ga ( SEQ ID No: 775)
158	gga gtt gtG gag agc tt ( SEQ ID No: 776)
159	cct aag agG gag tgt ca ( SEQ ID No: 777)
160	c ttc tat aaT cag gag gag ( SEQ ID No: 778)
161	ctg gac aga Cac ttc tat ( SEQ ID No: 779)
162	ag aag gac Ttc ctg gag ( SEQ ID No: 780)
163	cgg gcg gCg acg ga ( SEQ ID No: 781)
164	gc cag aag Aac atc ctg ( SEQ ID No: 782)
165	g gag ttc cAg gcg gtg ( SEQ ID No: 783)
166	caa gg gac Atc ctg gag c ( SEQ ID No: 784)
167	gac agg gCc gcc gc ( SEQ ID No: 785)
168	g cgg ttc cCg gac aga ( SEQ ID No: 786)
169	g gag ctg cGt aag tct g ( SEQ ID No: 787)
170	ctg gct tTc gct ggg g ( SEQ ID No: 788)
171	ttg gag ctg Tgt aag tct ( SEQ ID No: 789)
172	g gag ctg tGt aag tct g ( SEQ ID No: 790)
173	g tac ctg gaG aga tac tt ( SEQ ID No: 791)
174	cgg tac ctg Aac aga tac ( SEQ ID No: 792)
175	gag cag aAg cgg ggc ( SEQ ID No: 793)
176	g gag tac gCg cgc ttc ( SEQ ID No: 794)
177	ag ttc ctg Agc ttc gac ( SEQ ID No: 795)
178	cgt ttc ttg Gag ctg ctt ( SEQ ID No: 796)
179	ctg gag aga Cac ttc cat ( SEQ ID No: 797)
180	t tac tgc agG cac aac ta ( SEQ ID No: 798)

Table 22-7

Probe No.	Base Sequence
181	cct gat gcG gag tac tg ( SEQ ID No: 799)
182	g gag gag Aac gcg cg ( SEQ ID No: 800)
183	g gag aac gCg cgc ttc ( SEQ ID No: 801)
184	cgt ttc ttg Cag ctg ctt ( SEQ ID No: 802)
185	g gtg cgg Ctc ctg ga ( SEQ ID No: 803)
186	c ggg gtt gCt gag agc ( SEQ ID No: 804)
187	aac tac ggC gtt gtg ga ( SEQ ID No: 805)
188	g aca ttg aCg gtg ctg a ( SEQ ID No: 806)
189	c gag gtg gCc acc tac ( SEQ ID No: 807)
190	gtg tgg aaC ctg atc ag ( SEQ ID No: 808)
191	g gac acc taT tgc aga ta ( SEQ ID No: 809)
192	aac agt gaT ctg ggg ga ( SEQ ID No: 810)
193	tac tgc aga Tac aac tac g ( SEQ ID No: 811)
194	tgt cat ttc Ctc aat ggg ( SEQ ID No: 812)
195	ga gtg tgg Aac ctg atc ( SEQ ID No: 813)
196	c atg gca aAg ctg aca g ( SEQ ID No: 814)
197	cgt ttc ttg Cag cag gat ( SEQ ID No: 815)
198	ctg cac aga Ggc atc tat ( SEQ ID No: 816)
199	gaa gac aCg cgc gcc ( SEQ ID No: 817)
200	ac acg cgC gcc gcg ( SEQ ID No: 818)
201	c ctg gaa Aac agg cgc ( SEQ ID No: 819)
202	a ggt tcc tAc atg gca g ( SEQ ID No: 820)
203	tgt ttc ttg Cag cag gat ( SEQ ID No: 821)

Table 23-1

Allele Number		Probe Number for Detection							
DRB1*010101	0	2	3	4	5				
DRB1*010102	6								
DRB1*010201	7	8							
DRB1*010202	9								
DRB1*0103	10	11	12						
DRB1*0104	13	14							
DRB1*0105	15								
DRB1*0106	16	14							
DRB1*0107	17								
DRB1*0108	18								
DRB1*0109	19	16							
DRB1*0110	20								
DRB1*030101	21	22	23	24	25	26	27	14	
DRB1*030102	26	28	14						
DRB1*030201	29	30	31	23	24	26	27		
DRB1*030202	30	23	24	26	28				
DRB1*0303	30	31	23	24	26	27	14		
DRB1*0304	21	22	32	25	26	27	14		
DRB1*030501	21	22	23	24	25	26	27		
DRB1*030502	27	33							
DRB1*0306	21	34	22	23	24	26	27	14	
DRB1*0307	22	23	24	25	26	27	14		
DRB1*0308	23	35	36	26	27	14			
DRB1*0309	37								
DRB1*0310	38	26	27	14					
DRB1*0311	21	39	40	41	14				
DRB1*0312	42	26	27	14					
DRB1*0313	43	26	27	14					
DRB1*0314	21	22	23	24	25	26			
DRB1*0315	21	22	23	24	25	26	14		

Table 23-2

Allele Number		Probe Number for Detection							
DRB1*0316	44								
DRB1*0317	45	46	18	47	48				
DRB1*0318	49	14							
DRB1*0319	10	26	27	14					
DRB1*0320	27	8							
DRB1*0321	50	25	26	27	14				
DRB1*0322	51								
DRB1*0323	37	14							
DRB1*0324	25	39	40	48	14				
DRB1*0325	21	22	32	52	25	26	27	14	
DRB1*040101	53	20							
DRB1*040102	54								
DRB1*0402	53	12	14						
DRB1*040301	55	56	57	14					
DRB1*040302	55	58	57	14					
DRB1*0404	53	14							
DRB1*040501	55	59	60	56	61				
DRB1*040502	62								
DRB1*040503	63								
DRB1*040504	60	42	33						
DRB1*0406	55	60	57	14					
DRB1*040701	55	56	57						
DRB1*040702	64								
DRB1*0408	65	55	59	60	56				
DRB1*0409	60	61	20						
DRB1*0410	60	56	61	14					
DRB1*0411	53	57	14						
DRB1*0412	60	61	10	66	14				
DRB1*0413	60	20	14						
DRB1*0414	60	10	11	12					

Table 23-3

Allele Number		Probe Number for Detection				
DRB1*0415	55	36	67	68	14	
DRB1*0416	69					
DRB1*0417	60	61	57			
DRB1*0418	60	10	66	14		
DRB1*0419	65	55	59	60		
DRB1*0420	60	57				
DRB1*0421	60	20				
DRB1*0422	60	56	26	27	14	
DRB1*0423	70					
DRB1*0424	61	42	71			
DRB1*0425	60	56	67	66	14	
DRB1*0426	72					
DRB1*0427	56	57	8			
DRB1*0428	60	56	25	61		
DRB1*0429	73					
DRB1*0430	74					
DRB1*0431	55	60	56	75	76	
DRB1*0432	77					
DRB1*0433	78					
DRB1*0434	55	79	56	20		
DRB1*0435	55	25	20			
DRB1*0436	55	67	68	14		
DRB1*0437	55	80	81	14		
DRB1*0438	55	10	82			
DRB1*0439	83					
DRB1*0440	84					
DRB1*0441	55	85	86	57	14	
DRB1*0442	55	25	14			
DRB1*0443	55	60	25			
DRB1*0444	60	56	13	14		



Table 23-4

Allele Number		Probe Number for Detection				
DRB1*070101	87	88	89			
DRB1*070102	90	91	92	89		
DRB1*0703	93					
DRB1*0704	91	48				
DRB1*0705	94					
DRB1*0706	91	95	89			
DRB1*0707	96					
DRB1*080101	97	42	67	66	33	
DRB1*080102	98					
DRB1*080201	99	33				
DRB1*080202	97	18	67	66		
DRB1*080203	100					
DRB1*080302	45	97	61	10	66	
DRB1*080401	97	18	67	66	14	
DRB1*080402	18	67	66	101		
DRB1*080403	66	101	102			
DRB1*080404	66	14	103			
DRB1*0805	97	61	67	68		
DRB1*0806	61	67	66	14		
DRB1*0807	104	67	66	33		
DRB1*0808	38	105	66			
DRB1*0809	45	50	67	66	33	
DRB1*0810	97	61	10	66	14	
DRB1*0811	38	66	33			
DRB1*0812	10	66	8			
DRB1*0813	97	18	66	33		
DRB1*0814	106					
DRB1*0815	107	10	66			
DRB1*0816	108	33				
DRB1*0817	25	61	67	66		

Table 23-5

Allele Number		Probe Number for Detection					
DRB1*0818	45	97	61	10	109		
DRB1*0819	110	10	66				
DRB1*0820	111	18	67	66	14		
DRB1*0821	112						
DRB1*0822	8	113	114				
DRB1*0823	15	66					
DRB1*0824	97	18	67	68			
DRB1*090102	92	115					
DRB1*0902	58	115					
DRB1*100101	116						
DRB1*100102	117	118					
DRB1*110101	99	36	67	68			
DRB1*110102	36	67	68	33			
DRB1*110103	36	67	119	68	120		
DRB1*110104	121	18	25	35	67	68	
DRB1*1102	35	10	11	12	14		
DRB1*1103	99	122	14				
DRB1*110401	99	67	68	14			
DRB1*110402	36	14	103				
DRB1*1105	123	35	36	67	68		
DRB1*110601	36	67	68	8			
DRB1*110602	36	67	68	7	8		
DRB1*1107	35	36	26	27	14		
DRB1*110801	18	25	35	124			
DRB1*110802	36	124	33				
DRB1*1109	32	23	24	25	35	67	68
DRB1*1110	22	32	50	25	35	67	68
DRB1*1111	25	35	67	125	122		
DRB1*111201	126	25	35	67	68		
DRB1*111202	111	127	128	25	35	67	68

Table 23-6

Allele Number		Probe Number for Detection						
DRB1*1113	25	35	36	71	7	14		
DRB1*1114	35	10	11	12				
DRB1*1115	129	36	67	119	68			
DRB1*1116	23	35	10	11	12	14		
DRB1*1117	111	35	36	130	131	14		
DRB1*1118	18	35	10	109	14			
DRB1*1119	18	35	10	109				
DRB1*1120	23	35	10	11	12			
DRB1*1121	11	12	8					
DRB1*1122	55	25	36	67	68			
DRB1*1123	35	36	67	68	132	66		
DRB1*1124	108	36	67	119	68			
DRB1*1125	36	67	66	14				
DRB1*1126	133	134	18	25	35			
DRB1*112701	135	68	13					
DRB1*112702	35	68	136					
DRB1*1128	134	137	138	25	35	67	68	
DRB1*1129	45	111	134	25	35	67	68	
DRB1*1130	139	68						
DRB1*1131	35	140	10	109				
DRB1*1132	35	36	67	68	141			
DRB1*1133	142							
DRB1*1134	18	25	35	14				
DRB1*1135	142	14						
DRB1*1136	25	35	80	81	14			
DRB1*1137	45	111	134	18	35	67	68	
DRB1*1138	143							
DRB1*1139	144	68						
DRB1*1140	23	25	35	67	125	122	14	
DRB1*1141	35	67	125	122	14			

Table 23-7

Allele Number		Probe Number for Detection							
DRB1*1142	18	25	35	124	14				
DRB1*1143	144	68	14						
DRB1*120101	145	146	147	148	92	10	7	8	
DRB1*120102	145	146	147	148	92	10	8		
DRB1*120201	148	67	7	8					
DRB1*120202	148	67	120	8					
DRB1*120302	147	148	92	10	120				
DRB1*1204	148	36	10	7	8				
DRB1*1205	147	92	10	7	8				
DRB1*1206	147	148	92	10	7	8			
DRB1*1207	149								
DRB1*1208	150	148	92	10	7	8			
DRB1*130101	46	23	24	25	10	11	12	14	
DRB1*130102	151								
DRB1*130103	12	7	14						
DRB1*130201	46	23	24	25	10	11	12		
DRB1*130202	12	152							
DRB1*130301	42	109	153	33					
DRB1*130302	61	109	153						
DRB1*1304	25	61	11	12	14				
DRB1*1305	134	32	23	25	67	68			
DRB1*1306	46	23	25	10	109	14			
DRB1*130701	154	45	111	134	46	155	18	67	119
DRB1*130702	111	46	155	18	58	67	119	68	
DRB1*1308	46	50	11	12	14				
DRB1*1309	24	25	10	156	14				
DRB1*1310	46	23	25	10	109	153	14		
DRB1*1311	18	25	67	68	14				
DRB1*1312	111	61	10	109					

Table 23-8

Allele Number		Probe Number for Detection							
DRB1*1313	111	61	10	66					
DRB1*131401	18	25	67	119	68				
DRB1*131402	25	58	67	119	68				
DRB1*1315	30	25	11	12	14				
DRB1*1316	157								
DRB1*1317	97	12	14						
DRB1*1318	23	25	67	66	14				
DRB1*1319	30	50	11	12	14				
DRB1*1320	46	23	24	25	80	81	14		
DRB1*1321	111	25	61	67	68				
DRB1*1322	111	46	18	25	10	11	12	14	
DRB1*1323	11	12	33						
DRB1*1324	25	67	125	122	14				
DRB1*1325	154	45	111	134	46	18	25	124	
DRB1*1326	31	158	23	24	58	67	119	68	120
DRB1*1327	21	11	12	14					
DRB1*1328	159								
DRB1*1329	46	23	24	25	80	81			
DRB1*1330	25	61	10	109					
DRB1*1331	104	10	11	12					
DRB1*1332	23	61	11	12	14				
DRB1*1333	61	109	136						
DRB1*1334	160	11	12						
DRB1*1335	161								
DRB1*1336	46	23	24	10	11	12			
DRB1*1337	109	153	33						
DRB1*1338	61	11	12						
DRB1*1339	43	10	11	12					
DRB1*1340	46	23	24	10	11	12	14		

Table 23-9

Allele Number		Probe Number for Detection							
DRB1*1341	21	11	12						
DRB1*1342	23	67	68	14					
DRB1*1343	25	38	80	81	14				
DRB1*1344	111	134	46	18	25	14			
DRB1*1345	25	38	10	11	12				
DRB1*1346	18	104	162	67	135	68			
DRB1*1347	111	18	67	66	33				
DRB1*1348	61	11	12	14					
DRB1*1349	111	61	67	68					
DRB1*1350	134	137	25	67	68				
DRB1*1351	163								
DRB1*1352	46	32	52	25	10	11	12	14	
DRB1*1353	30	24	11	12	14				
DRB1*1354	92	125	122	14					
DRB1*1355	111	42	67	66	33				
DRB1*140101	99	111	130	131	14				
DRB1*140102	164	111	38	130	14				
DRB1*1402	99	158	23	24					
DRB1*1403	99	23	66						
DRB1*1404	99	97	130	131	14				
DRB1*140501	165	166	131	14					
DRB1*140502	165	131	14						
DRB1*1406	45	30	23	24	14				
DRB1*140701	164	111	38	130	131				
DRB1*140702	38	131	33						
DRB1*1408	164	111	107	130	131	14			
DRB1*1409	167	134	46	22	32	23			
DRB1*1410	59	38	130	131	14				
DRB1*1411	97	35	36	130	131	14			
DRB1*1412	30	23	24	66	14				

Table 23-10

Allele Number	Probe Number for Detection						
DRB1*1413	30	23	24	61			
DRB1*1414	111	50	130	131			
DRB1*1415	97	50	67	66	14		
DRB1*1416	38	10	11	12	14		
DRB1*1417	134	46	22	23	25	14	
DRB1*1418	23	24	166	130	131	14	
DRB1*1419	29	45	30	23	24	20	
DRB1*1420	133	150	30	50	14		
DRB1*1421	46	22	23	25	20	14	
DRB1*1422	50	38	105	67	135	68	
DRB1*1423	164	111	50	130	131	14	
DRB1*1424	30	158	23	24	10	168	156
DRB1*1425	111	18	38	105	67	135	68
DRB1*1426	169	14					
DRB1*1427	30	23	24	67	68	132	66
DRB1*1428	38	8	113				
DRB1*1429	30	158	23	24	8		
DRB1*1430	134	46	22	32	23	25	
DRB1*1431	97	38	7	14			
DRB1*1432	164	111	38	71	14		
DRB1*1433	24	25	57	14			
DRB1*1434	164	111	107	7	14		
DRB1*1435	25	38	130	131	14		
DRB1*1436	49	131					
DRB1*1437	165	156	14				
DRB1*1438	38	170	14				
DRB1*1439	171	38	130	131	14		
DRB1*1440	30	50	124	132	66		
DRB1*1441	45	111	150	30	50	172	
DRB1*1442	18	25	130	131			

Table 23-11

Allele Number		Probe Number for Detection						
DRB1*1443	173							
DRB1*1444	165	166	131					
DRB1*1445	165	10	131	14				
DRB1*150101	174							
DRB1*150102	175	176						
DRB1*150103	177	7	14					
DRB1*150104	177	25	10	156	14			
DRB1*150201	177	25	58	10	156			
DRB1*150202	25	10	168	156				
DRB1*150203	178							
DRB1*1503	177	179	25	58	10	156	14	
DRB1*1504	177	67	180	14				
DRB1*1505	177	25	58	16	14			
DRB1*1506	181							
DRB1*1507	177	58	10	156	14			
DRB1*1508	182							
DRB1*1509	183	156						
DRB1*1510	177	12	14					
DRB1*1511	177	58	10	156				
DRB1*1512	177	61	42	10	156	14		
DRB1*1513	177	25	58	184	156	14		
DRB1*160101	177	67	120					
DRB1*160102	177	67	68					
DRB1*160201	177	120						
DRB1*160202	177	124						
DRB1*1603	185							
DRB1*1604	127	58	67	68	132	66		
DRB1*1605	177	10	120					
DRB1*1607	186							
DRB1*1608	177	187	67	120				



Table 23-12

Allele Number		Probe Number for Detection					
DRB3*010101	188	34	172	162	26	28	
DRB3*01010201	189	26					
DRB3*010103	188	34	172	26	28		
DRB3*010104	28	175					
DRB3*0102	190	191	34	172	162	26	28
DRB3*0103	188	192	172	162	26	28	
DRB3*0104	193	34	172	162	26	28	
DRB3*0105	194	28					
DRB3*0106	188	34	50	162	26	28	
DRB3*0107	188	20	40	48			
DRB3*0108	188	23	24	162	26	28	
DRB3*0109	188	195	162	26	28		
DRB3*0110	196						
DRB3*0201	189	14					
DRB3*020201	197	198	195	47	48		
DRB3*020202	198	195	47	40	41		
DRB3*020203	199						
DRB3*020204	47	200	48				
DRB3*0203	198	201	47	48			
DRB3*0204	47	26	27	14			
DRB3*0205	30	195	47	48			
DRB3*0206	23	202	47	48			
DRB3*0207	47	104	162	48			
DRB3*0208	47	61	42	48			
DRB3*0209	195	92	40	48			
DRB3*0210	197	198	195	40	48		
DRB3*0211	47	10	48				
DRB3*0212	198	195	47	48			
DRB3*0213	203						
DRB3*0214	204						

Table 23-13

Allele Number		Probe Number for Detection					
DRB3*0215	198	195	47	40			
DRB3*0216	47	105	48				
DRB3*0217	47	67	48				
DRB3*030101	92	48	14				
DRB3*030102	205						
DRB3*0302	198	92	48	14			
DRB3*0303	30	50	162	92	26	28	
DRB4*010101	206						
DRB4*0102	207						
DRB4*010302	208	209	210				
DRB4*010303	206	131					
DRB4*010304	211						
DRB4*0104	212	213					
DRB4*0105	208	214					
DRB4*0106	208	209	210				
DRB4*0201N	87	14					
DRB5*010101	215						
DRB5*010102	129	58	67	119	68		
DRB5*0102	2	216	217	67	119	120	
DRB5*0103	218	219	220				
DRB5*0104	129	66					
DRB5*0105	108	67	119	120			
DRB5*0106	129	113					
DRB5*0107	129	10	221	120			
DRB5*0109	222						
DRB5*0110N	218	217	67	119	120		
DRB5*0111	129	156					
DRB5*0112	129	223	224	225			
DRB5*0202	226	113					
DRB5*0203	218	217	10	168	156		
DRB5*0204	218	67	180	113			
DRB5*0205	218	217	113				

Table 24-1

Allele Number		Probe Number for Detection						
DRB1*010101	0	1	2	3	4	5		
DRB1*010102	6							
DRB1*010201	7	8						
DRB1*010202	9							
DRB1*0103	10	11	12					
DRB1*0104	13	14						
DRB1*0105	15							
DRB1*0106	16	14						
DRB1*0107	17							
DRB1*0108	18							
DRB1*0109	19	16						
DRB1*0110	20							
DRB1*030101	21	22	23	14				
DRB1*030102	24	13	14					
DRB1*030201	21	25	23					
DRB1*030202	21	13						
DRB1*0303	25	26	27	28	24	23	14	
DRB1*0304	22	26	29	30	24	23	14	
DRB1*030501	22	26	27	28	30	24	23	
DRB1*030502	23	31						
DRB1*0306	22	32	26	27	28	24	23	14
DRB1*0307	21	23	14					
DRB1*0308	21	33	34	23	14			
DRB1*0309	35							
DRB1*0310	36	24	23	14				
DRB1*0311	22	37	38	23	14			
DRB1*0312	39	40	24	23				
DRB1*0313	41	24	23	14				
DRB1*0314	22	26	27	28	30	24		

Table 24-2

Allele Number	Probe Number for Detection							
DRB1*0315	22	26	27	28	30	24	14	
DRB1*0316	42							
DRB1*0317	43	44	18	45	13			
DRB1*0318	46	14						
DRB1*0319	47	24	23	14				
DRB1*0320	23	8						
DRB1*0321	48	30	24	23	14			
DRB1*0322	49							
DRB1*0323	35	14						
DRB1*0324	30	37	38	13	14			
DRB1*0325	22	26	50	51	30	24	23	14
DRB1*040101	52	20						
DRB1*040102	53							
DRB1*0402	52	12	14					
DRB1*040301	54	18	55	14				
DRB1*040302	54	56	55	14				
DRB1*0404	52	14						
DRB1*040501	54	57	58	18	39			
DRB1*040502	59							
DRB1*040503	54	57	58	18	39			
DRB1*040504	58	40	31					
DRB1*0406	54	58	55	14				
DRB1*040701	54	18	55					
DRB1*040702	60							
DRB1*0408	61	54	57	58	18			
DRB1*0409	58	39	20					
DRB1*0410	58	18	39	14				
DRB1*0411	52	55	14					
DRB1*0412	58	39	10	62	14			
DRB1*0413	58	20	14					

Table 24-3

Allele Number		Probe Number for Detection				
DRB1*0414	58	10	11	12		
DRB1*0415	54	58	34	63	64	
DRB1*0416	65					
DRB1*0417	58	39	55			
DRB1*0418	58	10	62	14		
DRB1*0419	61	54	57	58		
DRB1*0420	58	55				
DRB1*0421	61	54	57	20		
DRB1*0422	58	18	24	23	14	
DRB1*0423	66					
DRB1*0424	39	40	67			
DRB1*0425	58	18	63	64	68	62
DRB1*0426	69					
DRB1*0427	18	55	8			
DRB1*0428	58	18	30	39		
DRB1*0429	70					
DRB1*0430	71					
DRB1*0431	54	58	18	68	62	
DRB1*0432	72					
DRB1*0433	73					
DRB1*0434	74	75	18	20		
DRB1*0435	54	30	20			
DRB1*0436	54	63	64	14		
DRB1*0437	54	11	12	14		
DRB1*0438	54	47	20			
DRB1*0439	76					
DRB1*0440	77					
DRB1*0441	54	78	79	55	14	
DRB1*0442	54	30	14			
DRB1*0443	54	58	30			

Table 24-4

Allele Number	Probe Number for Detection				
DRB1*0444	58	18	13	14	
DRB1*070101	80	37	81		
DRB1*070102	82	83	84	81	
DRB1*0703	85				
DRB1*0704	83	13			
DRB1*0705	86				
DRB1*0706	83	87	81		
DRB1*0707	88				
DRB1*080101	89	40	63	62	31
DRB1*080102	90				
DRB1*080201	91	31			
DRB1*080202	89	18	63	62	
DRB1*080203	92				
DRB1*080302	21	10	62		
DRB1*080401	21	62	14		
DRB1*080402	18	63	62	93	
DRB1*080403	62	93	31		
DRB1*080404	62	14	31		
DRB1*0805	89	39	63	64	
DRB1*0806	39	63	62	14	
DRB1*0807	94	63	62	31	
DRB1*0808	36	95	62		
DRB1*0809	96	48	63	62	31
DRB1*0810	89	39	10	62	14
DRB1*0811	97	62			
DRB1*0812	10	62	8		
DRB1*0813	96	89	18	62	
DRB1*0814	98				
DRB1*0815	95	10	62		
DRB1*0816	99	31			

Table 24-5

Allele Number	Probe Number for Detection						
	30	39	63	62			
DRB1*0817	30	39	63	62			
DRB1*0818	96	89	39	10	64		
DRB1*0819	100	10	62				
DRB1*0820	101	18	63	62	14		
DRB1*0821	102						
DRB1*0822	8	103	31				
DRB1*0823	15	62					
DRB1*0824	89	18	63	64			
DRB1*090102	104	84					
DRB1*0902	104	56					
DRB1*100101	105						
DRB1*100102	106	107					
DRB1*110101	91	34	63	64			
DRB1*110102	34	63	64	31			
DRB1*110103	34	63	108	109	110		
DRB1*110104	111	18	30	33	63	64	
DRB1*1102	21	34	10	11	12	14	
DRB1*1103	91	12	14				
DRB1*110401	91	63	64	14			
DRB1*110402	34	14	31				
DRB1*1105	112	33	34	63	64		
DRB1*110601	34	63	64	8			
DRB1*110602	34	63	64	7	8		
DRB1*1107	33	34	24	23	14		
DRB1*110801	18	30	33	64			
DRB1*110802	18	30	33	64			
DRB1*1109	113	27	28	30	33	63	64
DRB1*1110	26	114	48	30	33	63	64
DRB1*1111	30	33	63	11	12		
DRB1*111201	115	30	33	63	64		

Table 24-6

Allele Number	Probe Number for Detection						
DRB1*111202	101	116	48	30	33	63	64
DRB1*11113	21	30	33	67	7	14	
DRB1*11114	21	34	10	11	12		
DRB1*11115	117	34	63	118	64		
DRB1*11116	27	33	10	11	12	14	
DRB1*11117	21	33	55	7	14		
DRB1*11118	18	33	10	64	14		
DRB1*11119	18	33	10	64			
DRB1*11120	27	33	10	11	12		
DRB1*11121	33	10	11	12			
DRB1*11122	54	30	34	63	64		
DRB1*11123	33	34	63	64	68	62	
DRB1*11124	99	34	63	118	64		
DRB1*11125	34	63	62	14			
DRB1*11126	43	101	119	18	30	33	
DRB1*112701	120	64	13				
DRB1*112702	33	64	23				
DRB1*11128	119	78	79	30	33	63	64
DRB1*11129	43	101	119	30	33	63	64
DRB1*11130	121	64					
DRB1*11131	122	123	10	64			
DRB1*11132	33	34	63	64	124		
DRB1*11133	125						
DRB1*11134	18	30	33	14			
DRB1*11135	125	14					
DRB1*11136	30	33	11	12	14		
DRB1*11137	43	101	119	18	33	63	64
DRB1*11138	126						
DRB1*11139	45	64					
DRB1*11140	27	30	33	63	11	12	



Table 24-7

Allele Number	Probe Number for Detection							
DRB1*1141	33	63	11	12	14			
DRB1*1142	18	30	33	64	14			
DRB1*1143	45	64	14					
DRB1*120101	127	21	128	129	84	10	7	8
DRB1*120102	127	21	128	129	84	10	8	
DRB1*120201	129	63	7	8				
DRB1*120202	129	63	110					
DRB1*120302	128	129	84	10	110			
DRB1*1204	129	34	10	7				
DRB1*1205	128	84	10	7	8			
DRB1*1206	21	128	129	84	10	7	8	
DRB1*1207	130							
DRB1*1208	131	129	84	10	7	8		
DRB1*130101	21	27	30	10	11	12	14	
DRB1*130102	132							
DRB1*130103	12	7	14					
DRB1*130201	21	27	30	10	11	12		
DRB1*130202	133	134						
DRB1*130301	40	135	136	31				
DRB1*130302	39	135	136					
DRB1*1304	21	40	10	11	12	14		
DRB1*1305	119	113	27	30	63	64		
DRB1*1306	44	27	30	10	64	14		
DRB1*130701	137	43	101	119	44	138	18	63
DRB1*130702	101	44	138	18	56	63	118	64
DRB1*1308	44	48	11	12	14			
DRB1*1309	28	30	47	16	14			
DRB1*1310	44	27	30	10	135	136	14	
DRB1*1311	18	30	63	64	14			

Table 24-8

Allele Number	Probe Number for Detection								
DRB1*1312	101	39	10	64					
DRB1*1313	101	39	10	62					
DRB1*131401	18	30	63	118	64				
DRB1*131402	30	56	63	118	64				
DRB1*1315	25	30	11	12	14				
DRB1*1316	139								
DRB1*1317	21	89	30	10	11	12	14		
DRB1*1318	27	30	63	62	14				
DRB1*1319	21	48	10	11	12	14			
DRB1*1320	44	27	28	30	11	12	14		
DRB1*1321	21	40	63	64					
DRB1*1322	101	44	18	30	10	11	12	14	
DRB1*1323	11	12	31						
DRB1*1324	30	63	11	12	14				
DRB1*1325	137	43	101	119	44	18	30	64	
DRB1*1326	26	113	27	28	56	63	108	109	110
DRB1*1327	22	11	12	14					
DRB1*1328	140								
DRB1*1329	44	27	28	30	11	12			
DRB1*1330	30	39	10	64					
DRB1*1331	141	10	11	12					
DRB1*1332	27	39	11	12	14				
DRB1*1333	39	135	23						
DRB1*1334	142	11	12						
DRB1*1335	143								
DRB1*1336	44	27	28	10	11	12			
DRB1*1337	135	136	31						
DRB1*1338	39	11	12						
DRB1*1339	41	10	11	12					

Table 24-9

Allele Number	Probe Number for Detection							
DRB1*1340	44	27	28	10	11	12	14	
DRB1*1341	22	11	12					
DRB1*1342	27	63	64	14				
DRB1*1343	30	36	11	12	14			
DRB1*1344	101	119	44	18	30	14		
DRB1*1345	30	36	10	11	12			
DRB1*1346	18	141	144	63	120	64		
DRB1*1347	101	18	63	62	31			
DRB1*1348	39	11	12	14				
DRB1*1349	101	39	63	64				
DRB1*1350	119	78	30	63	64			
DRB1*1351	145							
DRB1*1352	44	50	51	30	10	11	12	14
DRB1*1353	25	28	11	12	14			
DRB1*1354	84	11	12	14				
DRB1*1355	101	40	63	62	31			
DRB1*140101	91	101	55	7	14			
DRB1*140102	146	101	36	67	55			
DRB1*1402	91	27	28					
DRB1*1403	91	27	62					
DRB1*1404	91	89	55	7	14			
DRB1*140501	147	148	7	14				
DRB1*140502	147	7	14					
DRB1*1406	149	43	25	27	28	14		
DRB1*140701	146	101	36	55	7			
DRB1*140702	36	7	31					
DRB1*1408	146	101	95	55	7	14		
DRB1*1409	43	119	44	26	113	27		
DRB1*1410	57	36	55	7	14			
DRB1*1411	89	33	34	55	7			
DRB1*1412	25	27	28	64	68	62		

Table 24-10

Allele Number	Probe Number for Detection						
DRB1*1413	25	27	28	39			
DRB1*1414	146	101	48	55	7		
DRB1*1415	89	48	63	62	14		
DRB1*1416	48	36	10	11	12		
DRB1*1417	119	44	26	27	30	14	
DRB1*1418	27	28	148	55	7	14	
DRB1*1419	21	25	27	28	20		
DRB1*1420	43	101	131	25	48		
DRB1*1421	44	26	27	30	20		
DRB1*1422	48	36	95	63	120	64	
DRB1*1423	146	101	48	55	7	14	
DRB1*1424	25	113	27	28	47	19	16
DRB1*1425	101	18	36	95	63	120	64
DRB1*1426	150	14					
DRB1*1427	25	27	28	63	64	68	62
DRB1*1428	36	8	103				
DRB1*1429	25	113	27	28	8		
DRB1*1430	119	44	26	113	27	30	
DRB1*1431	89	36	7	14			
DRB1*1432	146	101	36	67	14		
DRB1*1433	28	30	55	14			
DRB1*1434	146	101	95	7	14		
DRB1*1435	30	36	55	7	14		
DRB1*1436	151	7					
DRB1*1437	147	16	14				
DRB1*1438	36	13	14				
DRB1*1439	152	36	55	7	14		
DRB1*1440	25	48	64	68	62		
DRB1*1441	43	101	131	25	153	154	
DRB1*1442	18	30	55	7			
DRB1*1443	155						

Table 24-11

Allele Number	Probe Number for Detection						
DRB1*1444	147	148	7				
DRB1*1445	147	47	7	14			
DRB1*150101	156						
DRB1*150102	157	158					
DRB1*150103	159	7	14				
DRB1*150104	159	30	47	16	14		
DRB1*150201	159	30	56	47	16		
DRB1*150202	30	47	19	16			
DRB1*150203	160						
DRB1*1503	159	161	30	56	47	16	14
DRB1*1504	159	162	16	14			
DRB1*1505	159	30	56	16	14		
DRB1*1506	163						
DRB1*1507	159	56	47	16			
DRB1*1508	164						
DRB1*1509	165	16					
DRB1*1510	159	12					
DRB1*1511	159	56	47	16			
DRB1*1512	159	39	40	47	16	14	
DRB1*1513	159	30	56	166	16	14	
DRB1*160101	159	63	110				
DRB1*160102	159	63	64				
DRB1*160201	159	110					
DRB1*160202	159	64					
DRB1*1603	167						
DRB1*1604	159	62					
DRB1*1605	159	10	110				
DRB1*1607	168						
DRB1*1608	159	28	63	110			
DRB3*010101	169	32	154	144	24	13	
DRB3*01010201	170	24					

Table 24-12

Allele Number	Probe Number for Detection						
DRB3*010103	169	32	154	24	13		
DRB3*010104	169	32	154	144	24	13	
DRB3*0102	171	172	32	154	144	24	13
DRB3*0103	169	173	154	144	24	13	
DRB3*0104	169	32	154	144	24	13	
DRB3*0105	174	13					
DRB3*0106	169	32	48	144	24	13	
DRB3*0107	169	175	38	13			
DRB3*0108	169	27	28	144	24	13	
DRB3*0109	169	176	144	24	13		
DRB3*0110	177						
DRB3*0201	170	14					
DRB3*020201	178	179	176	45	13		
DRB3*020202	178	179	176	45	38	23	
DRB3*020203	180						
DRB3*020204	45	181	13				
DRB3*0203	179	29	45	13			
DRB3*0204	45	24	23	14			
DRB3*0205	178	25	176	45	13		
DRB3*0206	182	183	45	13			
DRB3*0207	45	141	144	13			
DRB3*0208	45	39	40	13			
DRB3*0209	176	84	38	13			
DRB3*0210	178	179	176	38	13		
DRB3*0211	45	47	13				
DRB3*0212	184	13					
DRB3*0213	185						
DRB3*0214	186						
DRB3*0215	178	179	176	45	38		
DRB3*0216	45	95	13				
DRB3*0217	45	162	13				

Table 24-13

Allele Number	Probe Number for Detection					
DRB3*030101	84	13	14			
DRB3*030102	187					
DRB3*0302	179	48	84	175	38	13
DRB3*0303	25	48	144	84	24	13
DRB4*010101	188					
DRB4*0102	189					
DRB4*010302	80	190	14			
DRB4*010303	188	191				
DRB4*010304	192					
DRB4*0104	23	193				
DRB4*0105	194	195				
DRB4*0106	194	190	193			
DRB4*0201N	80	14				
DRB5*010101	196					
DRB5*010102	117	56	63	118	64	
DRB5*0102	197	78	63	108	110	
DRB5*0103	198	199	200			
DRB5*0104	117	62				
DRB5*0105	99	63	108	110		
DRB5*0106	117	103				
DRB5*0107	117	10	108	110		
DRB5*0109	201					
DRB5*0110N	197	78	63	108	110	
DRB5*0111	117	16				
DRB5*0112	117	84	67	81		
DRB5*0202	202	103				
DRB5*0203	198	78	47	19	16	
DRB5*0204	203	162	16	103		
DRB5*0205	203	78	103			

(Example 13)

Probes for identification of HLA-MICA allele

Extraction of DNA from 1 ml of human blood was performed using GFX Genomic Blood DNA Purification

5 Kit from Amersham Biosciences in the same manner as in Example 1.

Next, quantitative PCR was carried out in the same manner as in Example 1 except that probes in Tables 25-1 and 25-2 were used and 2  $\mu$ l of the mixed  
10 primers consisting of 1  $\mu$ l each of respective solutions of the following primers (10 pmol/ $\mu$ l) and 6  $\mu$ l of ultra pure water:

AGTGGAGCCAGTGGACCCAAGA (SEQ ID NO: 104)

TGATGTTTTCTTCTTACAACAAC (SEQ ID NO: 105)

15 After PCR amplification, referring to Amp Plot and Dissociation curves on a display of 5700 software, and to the allele-probe list 1 (Tables 27-1 and 27-2), it was identified as MICA\*00201.

(Example 14)

20 Extraction of DNA from 1 ml of human blood was performed in the same way as in Example 3. PCR of human HLA-MICA was then performed in the same manner as in Example 2 except that 3  $\mu$ l of the mixed primer consisting of 1  $\mu$ l each of the solutions containing  
25 the following sequences at 10 pmol/ $\mu$ l respectively, and 12  $\mu$ l of ultra pure water were used:



GTCTTCGTTATAACCTCACGGT (SEQ ID NO:106)

GCTCGTGAGCCTGCAGGTCCTG (SEQ ID NO:107)

AGTGGAGCCAGTGGACCCAAGA (SEQ ID NO:108)

At the same time, a DNA microarray was prepared  
5 to identify the allele in the specimen described  
above in the same manner as in Example 2, except that  
probes in the probe list of Table 26-1 were used to  
form the probe spots respectively.

Then, hybridization was performed using the  
10 above specimen and the prepared DNA microarray in the  
same manner as in Example 2. The DNA microarray was  
air-dried and the fluorometry measurement was  
conducted with GenePix4000B (Axon). Referring to the  
allele-probe correspondence list 2 (Tables 28-1 and  
15 28-2), it was identified as MICA\*00201.

#### Allele list

MICA\*001

gtcttcgttataaacctcacggtgctgtcctgggatggatctgtgcagtcagggtttctcactgaggtacatctgga  
20 tggtcagcccttcttgcgcigtgacaggcagaaatgcagggaagccccaggacagtgggcagaagatgtcctg  
ggaaataagacatgggacagagagaccagAgacttgacagggaacggaaggaccctcaggaigacccctggctcata  
tcaaggaccagaaagaaggcttgcatctccctccaggagattagggctgtgagatccatgaagacaacagcaccag  
gagctcccagcatcttactacgatggggagctcttctctcccaaacctggagactAaggaatggacaatgcc  
cagctccctccagagctcagaccttggccatgaacgtcaggaatttcttgaaggaagatgccatgaagaccaagacac  
25 actatcacgctatgcatgcagactgccctgcaggaactacggcgatacttaAaatccGgcgtagctctgaggagaac  
agtgccecccatggatgaatgtcacccgcagcgaggccctcagagggaacattaccgtgacatgcagggtctctggc  
ttctatcccttggaatatcacactgaGctggcgctcaggatggggatcttctttagccacgacacccagcagtgggggg

atgtccigcctigatgggaatggaacctaccagacctgggtggccaccaggatttgccaaggagaggagcagaggtt  
caccitgctacatggaacacacgcggaatcacagcactcaccctgtgcccctcgggaaagtgctgggtgcttcagagt  
cattggcagacatccatgttctgctgttgctgctNNNgctgctNNNNNNNNNNNNNNNNNNat t t t t g t t a t t a  
t t a t t t c t a t g t c c g t t g t t g t a a g a g a a a c a t c a g c t g c a g a g g g t c c a g a g c t c g t g a g c c t g c a g g t c c t  
5 g g a t c a a c a c c c a g t t g g g a c g a g t g a c c a c a g g g a t g c c a c a c a g c t c g g a t t c a g c c t c t g a t g t c a g a t c t t  
g g g t c c a c t g g c t c c a c t (SEQ ID NO:109)

gagc|cccagcat|t|t|a|c|acgat|ggggagc|t|t|c|c|t|c|c|c|a|a|a|a|c|c|t|g|g|a|g|a|c|t|g|g|a|a|t|g|g|c|c|c|  
c|a|g|t|c|c|t|c|c|a|g|a|g|c|t|c|a|g|a|c|t|t|g|g|c|c|a|t|g|a|a|c|g|t|c|a|g|g|a|a|t|t|c|t|t|g|a|a|g|g|a|a|g|a|t|g|c|c|a|t|g|a|a|g|a|c|c|a|a|g|a|c|a|c|  
a|c|t|a|t|c|a|c|g|c|t|a|t|g|c|a|t|g|c|a|g|a|c|t|g|c|c|t|g|c|a|g|g|a|a|c|t|a|c|g|g|c|g|a|t|a|t|c|t|a|a|a|a|t|c|c|g|g|c|g|t|a|g|t|c|t|g|a|g|g|a|g|a|a|c|  
a|g|t|g|c|c|c|c|c|c|a|t|g|g|t|g|a|a|t|g|t|c|a|c|c|c|g|c|a|g|t|g|a|g|g|c|c|t|c|a|g|a|g|g|g|c|a|a|c|a|t|t|a|c|c|g|t|g|a|c|a|t|g|c|a|g|g|g|c|t|t|c|t|g|g|c|  
5 t|t|c|t|a|t|c|c|t|g|g|a|a|t|a|t|c|a|c|a|c|t|g|a|g|c|t|g|g|c|g|t|c|a|g|g|a|t|g|g|g|l|a|t|c|t|t|t|g|a|g|c|c|a|c|g|a|c|c|c|c|a|g|c|a|g|t|g|g|g|g|g|  
a|t|g|t|c|c|t|g|c|c|t|g|a|t|g|g|g|a|t|g|g|a|a|c|t|a|c|c|a|g|a|c|c|t|g|g|g|t|g|g|c|c|a|c|c|a|g|g|a|t|t|t|g|c|c|a|a|g|g|a|g|g|a|g|g|a|g|c|a|g|a|g|t|t|  
c|a|c|c|t|g|c|t|a|c|a|t|g|g|a|a|c|a|c|a|g|c|g|g|g|a|a|t|c|a|c|a|g|c|a|c|t|c|a|c|c|t|g|t|g|c|c|c|t|c|t|g|(SEQ ID NO:111)

MICA\*004

g l c t t c g t t a t a a c c t c a c g g t g c t g t c c t g g g a t g g a t c t g t g c a g t c a g g g t t t c t t g c t g a g g t a c a t c t g g a  
10 t g g t c a g c c c t t c c t g c g c t a t g a c a g g c a g a a a t g c a g g g c a a a g c c c c a g g g a c a g t g g g c a g a a g a t g t c c t g  
g g a a a t a a g a c a t g g g a c a g a g a g a c c a g g g a c t t g a c a g g g a a c g g a a g g a c c t c a g g a t g a c c c t g g c t c a t a  
t c a a g g a c c a g a a a g a a g g c t t g c a t t c c c t c c a g g a g a t t a g g g t c t g t g a g a t c c a t g a a g a c a a c a g c a c c a g  
g a g c t c c c a g c a t t t c t a c t a c g a t g g g g a g c t c t t c c t c t c c c a a a c g t g g a g a c t g a g g a a t g g a c a g t g c c c  
c a g t c c t c c a g a g c t c a g a c c t t g g c c a t g a a c g t c a g g a a t t t c t t g a a g g a a g a t g c c a t g a a g a c c a a g a c a c  
15 a c t a t c a c g c t a t g c a t g c a g a c t g c c t g c a g g a a c t a c g g c g a t a t c t l a g a a t c c a g c g t a g t c c t g a g g a g a a G  
a g t g c c c c c c a t g g t g a a t g t c a c c c g c a g c g a g g c c t c a g a g g g c a a c a t c a c c g t g a c a t g c a g g g c t t c c a g c  
t t c t a t c c c c g g a a t a t c a c a c t g a c c t g g c g t c a g g a t g g g g t a t c t t t g a g c c a c g a c a c c c a g c a g t g g g g g g  
a t g t c c t g c c t g a t g g g a a t g g a a c c t a c c a g a c c t g g g t g g c c a c c a g g a t t t g c c a a g g a g a g g a g c a g a g g t t  
c a c c t g c t a c a t g g a a c a c a g c g g g a a t c a c a g c a c t c a c c c t g t g c c c t c t g g g a a a g t g c t g g t g c t t c a g a g t  
20 c a t t g g c a g a c a t t c c a t g t t t c t g c t g t t g c t g c t g c t g c t g c t a t t t t t g t t a t t a t t a t t t t c t a t g t c c  
g t t g t t g l a a g a g a a a a c a t c a g c t g c a g a g g g t c c a g a g c t c g t g a g c c t g c a g g t c c t g g a t c a a c a c c c a g t  
t g g g a c a g a t g a c c a c a g g g a t g c c a c a c a g c t c g g a t t t c a g c c t c t g a t g t c a g C t c t t g g g t c c a c t g g c t c c  
a c t (SEQ ID NO:112)

MICA\*005

25 gtccttcgttataaccctacaggctgcgtctctgggatggatctgtgcagtcagggttctctgctgagglacatctgga  
tggtcagcccttcttgcgctatgacaggcagaaaatgcaggggcaaagccccagggacagtgggcagaagaatgctctg  
ggaaaataagacatgggacagagagaccagggacttgacagggaacggaaaggacctcaggatgaccttggctcata



ggaaataagacatgggacagagagaccagggacttgacagggaaacggaaaggacctcaggatgacccitggctcata  
tcaaggaccagaaagaaggcttgcattccctccaggagattagggctgtgagatccatgaagacaacagcaccag  
gagctcccagcatttctactacgatggggagctcttctctcccaaaacctggagactgaggaatggacaatgccc  
cagctctccagagctcagaccttggccatgaacgtcaggaatttcttgaaggaagatgccatgaagaccaagacac  
5 actatcacgctatgcatgcagactgccitgcaggaactacggcgatactctaaaatccggcgtagtccitaggagaaac  
agtgccccccatggatgaatgtcacccgcagTgaggcctcagagggaacattaccgtgacatgcagggttcttggc  
ttctatcccttggaatatcacactgagctggcgctcaggatggggatcttctttagccacgacaccagcagtgggggg  
atgtcttgcctgatgggaatggaacctaccagacctgggtggccaccaggatttgccaaggagaggagcagaggtt  
cacctgctacatggaacacagcgggaatcacagcactcaccttctgccccttgggaaagtgttggtgcttcagagt  
10 cattggcagacattccatgtttctgctgttgcgtgctgctatcttctgttattattatttctatgtccgttgtt  
gtaagaagaaaacatcagctcagaggggtccag (SEQ ID NO:115)

MICA\*00702

gtcttcgttataacctcacggtgctgttcttgggatggatctgtgcagtcagggtttcttcgtgaggatcatctgga  
tggtcagcccttcttgcgctatgacaggcagaaatgcagggcaaagccccaggacagtgggcagaagatgtcctg  
15 ggaaataagacatgggacagagagaccagGgacttgacagggaaacggaaaggacctcaggatgacccitggctcata  
tcaaggaccagaaagaaggcttgcattccctccaggagattagggctgtgagatccatgaagacaacagcaccag  
gagctcccagcatttctactacgatggggagctcttctctcccaaaacctggagactGaggaatggacaatgccc  
cagctctccagagctcagaccttggccatgaacgtcaggaatttcttgaaggaagatgccatgaagaccaagacac  
actatcacgctatgcatgcagactgccitgcaggaactacggcgatactctaaaatccggcgtagtccitaggagaaac  
20 agtgccccccatggatgaatgtcacccgcagcaggcctcagagggaacattaccgtgacatgcagggttcttggc  
ttctatcccttggaatatcacactgagctggcgctcaggatggggatcttctttagccacgacaccagcagtgggggg  
atgtcttgcctgatgggaatggaacctaccagacctgggtggccaccaggatttgccaaggagaggagcagaggtt  
cacctgctacatggaacacagcgggaatcacagcactcaccttctgcccctctg (SEQ ID NO:116)

MICA\*00801

gtcttcgttataacctcacggtgctgttcttgggatggatctgtgcagtcagggtttcttgcitaggatcatctgga  
tggtcagcccttcttgcgctatgacaggcagaaatgcagggcaaagccccaggacagtgggcagaagatgtcctg  
25 ggaaataagacatgggacagagagaccagggacttgacagggaaacggaaaggacctcaggatgacccitggctcata

[illegible]

MI CA\*00802

gtcttcgttataaaccacacgggtgcgtgctcctgggaaggatctgtgcagtcagggttctctgctgaggtacatctggga  
15 tggtcagcccttccctgcgcctatgacaggcagaaaaatgcagggcacaagccccagggacagctgggcagaagaatgtccttg  
ggaaataagacatgggacagagagaccagggacttgacagggaacggaaaggaccacaggatgaccttggctcata  
tcaaggaccagaaagaaggcttgcattccctccaggagattagggctctgtgagatccatgaagacaacagcaccag  
gagctcccagcatttctactacgatggggagctcttccctctcccaaaacctggagactgaggaatggacagtgtccc  
cagctccctccagagctcagaccttggccatgaacgtcaggaattcttgaaggaagatgccatgaagaccaagacac  
20 acatcacgcctatgcatgcagactgcttgcaggaactacggcgatactagaatccggcgtagtccctgaggagaac  
agtgtccccccatgggtgaatgtcacccgcagcgaggcctcagagggcaacatcaccttgacatgcagggcttccagc  
ttctatccccggaalatcatactgacctggcgctcaggatggggatctctttagccacgacacccagcagctgggggg  
atgtccttgcctgatgggaatggaacctaccagacctgggttggccacTaggaatttggcgaggagaggagcagaggtt  
caccctgctacatggaacacagcgggaatcacagcactcaccttctgcccctctgggaaagtgtgtgtgtcttcagagt  
25 cattggcagacattccatgtttctgctgtttgctgtgtggctgctgtcTatttttgttattattattttctatgtccgt  
tgttgttaagaagaaaacatcagctgcagaggggtccag (SEQ ID NO:118)

MICA\*00803

gtcttcgttataaacctcacgggtgcgtgctctgggaatggatctgtgcagtcagggtttcttgc t gaggtaacatctggga  
tggtcagcccttcttgcgc t atgacaggcagaaaatgcagggc aaagccccagggaacag tgggcagaaga t g t c c t g  
ggaaa taagaca tgggacagagagaccaggga c t gacagggaacggaaaggacctcaggatgacctgggtcata  
tcaaggaccagaaagaaggcttgcat tccctccaggagat t agggctgtgagatccatgaagacaacagcaccag  
5 gaggctcccagcatttctactacgatggggagctcttctctcccaaaacctggagac t gaggaa tggacagt gccc  
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a c t a t c a c g c t a t g c a t g c a g a c t g c c t g c a g g a a c t a c g g c g a t a t c t a g a a t c c g g c g t a g t c c t g a g g a g a a c  
a g t g c c c c c c a t g g t g a a t g t c a c c c g c a g c g a g g c A t c a g a g g g c a a c a t c a c c g t g a c a t g c a g g g c t t c c a g c  
t t c t a t c c c c g g a a t a t c a t a c t g a c c t g g c g t c a g g a t g g g g t a t c t t t g a g c c a c g a c a c c a g c a g t g g g g g g  
10 a t g t c c t g c c t g a t g g g a a t g g a a c c t a c c a g a c c t g g g t g g c c a c c a g g a t t t g c c a g g a g a g g a g c a g a g g t t  
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MICA\*010

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MICA\*011

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MICA\*014

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20 MICA\*015

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MICA\*016

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MICA\*017

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[illegible]

[illegible]

## MICA\*021

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## MICA\*022

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## 25 MICA\*023

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MICA\*030

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MICA\*031

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agtgccccccatggatgaatgtcacccgcagcgaggccctcagagggcaacattaccgtgacatgcagggcttctggc  
5 tttatccctggaaatcacactgagctggcgctcaggatggggatctttagccacgacaccagcagtgggggg  
atgtcttgcttgatgggaatggaacctaccagacctgggtggccaccaggatttgccaaggagaggagcagaggtt  
cacctgctacatggaacacagcgggaatcacagcacacacctgtgccctctg (SEQ ID NO:144)

MICA\*032

gtcttcgttataaaccacaggctgtcttgggatggatctgtgcagtcagggttcttgcctgaggtacatctgga  
10 tggtcagcccttcttgcgtatgacaggcagaaaatgcagggcaaagccccaggacagtgggcagaagatgtcttg  
ggaaataagacatgggacagagagaccagggaacttgacagggaacggaaaggacctcaggatgaccttggctcata  
tcaaggaccagaaagaaggcttgcatctccctccaggagattagggtctgtgagatccatgaagacaacagcaccag  
gagctcccagcatcttctactacgatggggagctcttcccttcccaaaaccaggagactaggaatggacaatgccc  
cagctccctccagagctcagaccttggccaatgaacgtcaggaatttcttgaaggaagatgccatgaagaccaagacac  
15 TctatcacgctatgcatgcagactgccatgcaggaactacggcgataatctaaaatccAgcgtagctctgaggagaaG  
agtgtccccccatggatgaatgtcacccgcagcgaggccctcagagggcaacattaccgtgacatgcagggcttctggc  
ttctatccctggaaatcacactgagctggcgctcaggatggggatctttagccacgacaccagcagtgggggg  
atgtcttgcttgatgggaatggaacctaccagacctgggtggccaccaggatttgccaaggagaggagcagaggtt  
cacctgctacatggaacacagcgggaatcacagcacacacctgtgccctctg (SEQ ID NO:145)

20 MICA\*033

gtcttcgttataaaccacaggctgtcttgggatggatctgtgcagtcagggttcttgcctgaggtacatctgga  
tggtcagcccttcttgcgtatgacaggcagaaaatgcagggcaaagccccaggacagtgggcagaagatgtcttg  
ggaaataagacatgggacagagagaccagggaacttgacagggaacggaaaggacctcaggatgaccttggctcata  
tcaaggaccagaaagaaggcttgcatctccctccaggagattagggtctgtgagatccatgaagacaacagcaccag  
25 gagctcccagcatcttctactacgatggggagctcttcccttcccaaaaccaggagTcttaggaatggacagtggcc  
cagctccctccagagctcagaccttggccaatgaacgtcaggaatttcttgaaggaagatgccatgaagaccaagacac  
actatcacgctatgcatgcagactgccatgcaggaactacggcgataatctagaatccagcgtagctctgaggagaac

[illegible]

MICA\*034

giccttcgtatataaccctcacggctgcctccggggaatggatctgtgcagtcagggtttctcgcctgaggatcatcttgga  
tggtcagcccttctctgcctgtgacaggcagaaaatgcaggggcaaagccccagggaacagtgggcagaagaatgtccctg  
10 ggaaaataagacatgggacagagagaccagggaacttgacagggaacggaaaggacctcaggatgacctgggtcata  
tcaaggaccagaaaagaaggcttgcattccctccaggagattagggctctgtgagatccatgaagacaacagcaccag  
gagctcccagcatcttactacgatggggagctcttctctcccaaaacctggagactgaggaatggacaatgccc  
cagctccctccagagctcagaccttggccatgaacgtcaggaatttcttgaaggaagatgccGigaagaccaagacac  
actatcacgctatgcatgcagactgcttgcaggaactacggcgatacttaaaatccggcgtagtccctgaggagaac  
15 agtgccecccatggatgaatgtcacccgcagcgaggccctcagagggaacatcacccctgacatgcagggtctccagc  
ttctatccccggaatatcacactgaCctggcgctcaggatggggatcttttagccacgacaccagcagatgggggg  
atgtctctgctgaatgggaatggaacctaccagacctgggtggccaccaggatttgccaaggagaggagcagaggtt  
caccctgctacatggaacacagcgggaatcacagcactcaccctgtgccctctg(SAQ ID NO:147)

MICA\*035

20 gtcttcgttataacctcacggctgcctccGgggatggatctgtgcagtcagggcttcctcgctgagggtacatctggga  
tggtcagcccttccctgcgcctgtgacaggcagaaaatgcagggcacaagccccagggacagtgggcagaagatgtccctg  
ggaaaataagacatgggacagagagaccagggaacttgacagggaacggaaaggacctcaggatgacctggcctcata  
tcaaggaccagaaagaaggcttgcatctccctccaggagattagggtctgtgagatccatgaagacaacagcaccag  
gagctcccagcatcttctacacgatggggagctcttccctctcccaaaacctgggagactgaggaatggacaatgccc  
25 cagtcctccagagctcagaccttggccatgaacgtcaggaattcttgaaggaagatgccatgaagaccaagacac  
actatcacgctatgcatgcagactgcctgcaggaactacggcgatatctaaaaatccggcgtagtcttgaggagaaac  
agtgcaccccccatggigaatgtcaccgcagcgaggcctcagagggcacaatcacccgtgacatgcagggcttccagc

t t c t a t c c c c g g a a t a t c a T a c t g a c c t g g c g t c a g g a t g g g g t a t c t t t g a g c c a c g a c a c c c a g c a g t g g g g g g  
a t g t c c t g c c t g a t g g g a a t g g a a c c t a c c a g a c c t g g g t g g c c a c c a g g a t t t g c c g a g g a g a g g a g c a g a g g t t  
c a c c t g c t a c a t g g a a c a c a g c g g g a a t c a c a g c a c t c a c c c t g t g c c c t c t g (SEQ ID NO:148)

MICA\*036

5 g t c t t c g t t a a a c c t c a c g g t g c t g t c c g g g a t g g a t c t g t g c a g t c a g g g t t t c t c g c t g a g g t a c a t c t g g a  
t g g t c a g c c c t t c c t g c g c t g t g a c a g g c a g a a a t g c a g g g c a a a g c c c c a g g g a c a g t g g g c a g a a g a t g t c c t g  
g g a a a t a a g a c a t g g g a c a g a g a g a c c a g g g a c t t g a c a g g g a a c g g a a g g a c c t c a g g a t g a c c c t g g c t c a t a  
t c a a g g a c c a g a a a g a g g c t t g c a t t c c c t c c a g g a g a t t a g g g t c t g t g a g a t c c a t g a a g a c a a c a g c a c c a A  
g a g c t c c c a g c a t t t c t a c t a c g a t g g g g a g c t c t t c c t c t c c a a a a c c t g g a g a c t g a g g a a t g g a c a a t g c c c  
10 c a g t c c t c c a g a g c t c a g a c c t t g g c c a t g a a c g t c a g g a a t t t c t t g a a g g a a g a t g c c a t g a a g a c c a a g a c a c  
a c t a t c a c g c t a t g c a t g c a g a c t g c c t g c a g g a a c t a c g g c g a t a t c t a g a a t c c a g c g t a g t c c t g a g g a g a a c  
a g t g c c c c c a t g g t g a a t g t c a c c c g c a g c g a g g c c t c a g a g g g c a a c a t t a c c g t g a c a t g c a g g g c t t c t g g c  
t t c t a t c c c t g g a a t a t c a c a c t g a g c t g g c g t c a g g a t g g g g t a t c t t t g a g c c a c g a c a c c c a g c a g t g g g g g g  
a t g t c c t g c c t g a t g g g a a t g g a a c c t a c c a g a c c t g g g t g g c c a c c a g g a t t t g c c a a g g a g a g g a g c a g a g g t t  
15 c a c c t g c t a c a t g g a a c a c a g c g g g a a t c a c a g c a c t c a c c c t g t g c c c t c t g (SEQ ID NO:149)

MICA\*037

g t c t t c g t t a a a c c t c a c g g t g c t g t c c t g g g a t g g a t c t g t g c a g t c a g g g t t t c t c g c t g a g g t a c a t c t g g a  
t g g t c a g c c c t t c c t g c g c t g t g a c a g g c a g a a a t g c a g g g c a a a g c c c c a g g g a c a g t g g g c a g a a g a t g t c c t g  
g g a a a t a a g a c a t g g g a c a g a g a g a c c a g g g a c t t g a c a g g g a a c g g a a g g a c c t c a g g a t g a c c c t g g c t c a t a  
20 t c a a g g a c c a g a a a g a g g c t t g c a t t c c c t c c a g g a g a t t a g g g t c t g t g a g a t c c a t g a a g a c a a c a g c a c c a g  
g a g c t c c c a g c a t t t c t a c t a c g a t g g g g a g c t c t t c c t c t c c a a a a c c t g g a g a c t g a g g a a t g g a c a a t g c c c  
c a g t c c t c c a g a g c t c a g a c c t t g g c c a t g a a c g t c a g g a a t t t c t t g a a g g a a g a t g c c a t g a a g a c c a a g a c a c  
a c t a t c a c g c t a t g c a t g c a g a c t g c c t g c a g g a a c t a c g g c g a t a t c t a a a a t c c g g c g t a g t c c t g a g g a g a a c  
a g t g c c c c c a t g g t g a a t g t c a c c c g c a g c g a g g c c t c a g a g g g c a a c a t c a c c g t g a c a t g c a g g g c t t c c a g c  
25 t t c t a t c c c c g g a a t a t c a T a c t g a c c t g g c g t c a g g a t g g g g t a t c t t t g a g c c a c g a c a c c c a g c a g t g g g g g g  
a t g t c c t g c c t g a t g g g a a t g g a a c c t a c c a g a c c t g g g t g g c c a c c a g g a t t t g c c g a g g a g a g g a g c a g a g g t t  
c a c c t g c t a c a t g g a a c a c a g c g g g a a t c a c a g c a c t c a c c c t g t g c c c t c t g (SEQ ID NO:150)

MICA\*038

gtcttcgttataacctcacggctgcttcctgggatggatctgtgcagtcagggtttctcgctgaggtacatctgga  
tggtcagcccttccctgcgcctgacaggcagaaatgcagggcaaagcccaggacagtgggcagaagaatgtccctg  
ggaaataagacatgggacagagagaccagggaactgacagggaacggaaaggacctcaggatgaccttggtcata  
5 tcaaggaccagaaagaaggcttgcatccctccaggagattagggtctgtgagatccatgaagacaacagcaccag  
gagctcccagcatttctactacgatggggagctcttccctcccaaaacctggagactgaggaatggacaatgccc  
cagctctccagagctcagaccttgccatgaacgtcaggaatttctgaaggaagatgccatgaagaccaagacac  
actatcacgctatgcatgcagactgccctgcaggaactacggcgatatctaaaaaccggcgtagtcttgaggagaac  
agtgtcccccatggatgaatgtcacccgcagcaggccctcagagggaacaatCaccgtgacatgcagggttccagc  
10 ttctatccccggaatatcacactgacctggcgctcaggatgggtatctttgagccacgacaccagcagtgggggg  
atgtccctgcctgatgggaatggaacctaccagacctgggtggccaccaggatttgccaaggagaggagcagaggtt  
caccctgctacatggaacacacgggaatcacagcactcaccctgtgccccttg (SEQ ID NO:151)

MICA\*039

gtcttcgttataacctcacggctgcttcctgggatggatctgtgcagtcagggtttctcgctgaggtacatctgga  
15 tggtcagcccttccctgcgcctgacaggcagaaatgcagggcaaagcccaggacagtgggcagaagaatgtccctg  
ggaaataagacatgggacagagagaccagggaactgacagggaacggaaaggacctcaggatgaccttggtcata  
tcaaggaccagaaagaaggcttgcatccctccaggagattagggtctgtgagatccatgaagacaacagcaccag  
gagctcccagcatttctactacgatggggagctcttccctcccaaaacctggagactgaggaatggacaatgccc  
cagctctccagagctcagaccttgccatgaacgtcaggaatttctgaaggaagatgccatgaagaccaagacac  
20 actatcacgctatgcatgcagactgccctgcaggaactacggcgatatctaaaaaccggcgtagtcttgaggagaac  
agtgtcccccatggatgaatgtcacccgcagcaggccctcagagggaacaatcaccgtgacatgcagggttccagc  
ttctatccccggaatatcatctgacctggcgctcaggatgggtatctttgagccacgacaccagcagtgggggg  
atgtccctgcctgatgggaatggaacctaccagacctgggtggccaccaggatttgccgaggagaggagcagaggtt  
caccctgctacatggaacacacgggaatcacagcactcaccctgtgccccttg (SEQ ID NO:152)

25 MICA\*040

gtcttcgttataacctcacggctgcttcctgggatggatctgtgcagtcagggtttctcgctgaggtacatctgga  
tggtcagcccttccctgcgcctgacaggcagaaatgcagggcaaagcccaggacagtgggcagaagaatgtccctg







gagctcccagcatcttctactacgatggggagctcttccctctcccaaaacgtggagactgaggaatggacagtgtccc  
cagtcctccagagctcagaccttggccaatgaacgtcaggaatttcttgaaggaagatgccatgaagaccaagacac  
actatcagcctatgcatgcagactgacctgcaggaactacggcgatactagaatccagcgtagctctgaggagaaG  
agtgtcccccaatggatgaatgtcaccgcagcgaggccctcagagggaacaatcacctgtacatgcagggtctccagc  
5 tttctatccccggaatatcacactgaCctggcgctcaggatggggatcttcttgagccacgacacccagcagtgggggg  
atgtcctgtcctgatgggaatggaacctaccagacctgggtggccaccaggatttgcgaaggagaggagcagagggt  
cacctgtctacatggaacacagcggaatcacagcactcacctgtgcccctctg (SEQ ID NO:157)

MICA\*045

gtcttcgttataaacctcacggctgtcttgggaatggatctgtgcagtcagggtttctcgcctgaggtacatctgga  
10 tggtcagcccttccctgcgcctgtgacaggcagaaatgcagggaagccccaggacagtgggcagaagatgtcctg  
ggaaataagacatgggacagagagaccagggaacttgacagggaacggaaaggacctcaggatgacctggctcata  
tcaaggaccagaaagaaggcttgcatctccctccaggagattagggtctgtgagatccatgaagacaacagcaccag  
gagctcccagcatcttctactacgatggggagctcttccctctcccaaaacctggagactgaggaatggacaatgtccc  
cagtcctccagagctcagaccttggccaatgaacgtcaggaatttcttgaaggaagatgccatgaagaccaagacac  
15 actatcagcctatgcatgcagactgacctgcaggaactacggcgatacttaaaatccggcgtagctctgaggagaa  
agtgtcccccaatggatgaatgtcaccgcagtcaggccctcagagggaacaattaccgtgacatgcagggtctctggc  
ttctatcccgggaatatcacactgagctggcgctcaggatggggatcttcttgagccacgacacccagcagtgggggg  
atgtcctgtcctgatgggaatggaacctaccagacctgggtggccaccaggatttgcGaaggagaggagcagagggt  
cacctgtctacatggaacacagcggaatcacagcactcacctgtgcccctctgggaaagtgtgtgtcttcagagt  
20 catgtggcagacattccatgttctgtctgtgtgtgtgtctgtatcttctgttattattatttctatgtccgttgtt  
gtaagaagaaaacatcagctgcagagggtccag (SEQ ID NO:158)

MICA\*046

gtcttcgttataaacctcacggctgttccggggatggatctgtgcagtcagggtttctcgcctgaggtacatctgga  
tggtcagcccttccctgcgcctgtgacaggcagaaatgcagggaagccccaggacagtgggcagaagatgtcctg  
25 ggaaataagacatgggacagagagaccagggaacttgacagggaacggaaaggacctcaggatgacctggctcata  
tcaaggaccagaaagaaggcttgcatctccctccaggagattagggtctgtgagatccatgaagacaacagcaccag  
gagctcccagcatcttctactacgatggggagctcttccctctcccaaaacctggagactgaggaatggacaatgtccc





In the following, Probe List M1 and M2 are shown in Tables 25-1 and 25-2 and Tables 26-1 and 26-2 and Tables 27-1 and 27-2 and Tables 28-1 and 28-2 respectively.

5

Table 25-1

Probe No.	Base Sequence
0	tgg gac aga gag acc agA ( SEQ ID No: 1)
1	tcc caa aac ctg gag act A ( SEQ ID No: 2)
2	g gaa cta cgg cga tat cta A ( SEQ ID No: 3)
3	cgg cga tat cta aaa tcc G ( SEQ ID No: 4)
4	cc tgg aat atc aca ctg aG ( SEQ ID No: 5)
5	t att ttt gtt att att att ttc taC ( SEQ ID No: 6)
6	c ctc acg gtg ctg tcc G ( SEQ ID No: 7)
7	gtg aat gtc acc cgc agT ( SEQ ID No: 8)
8	c gta gtc ctg agg aga aG ( SEQ ID No: 9)
9	t cag cct ctg atg tca gC ( SEQ ID No: 10)
10	cag ccc ttc ctg cgc tA ( SEQ ID No: 11)
11	gag act gag gaa tgg aca G ( SEQ ID No: 12)
12	cc cgg aat atc aca ctg aC ( SEQ ID No: 13)
13	gcc acc agg att tgc cG ( SEQ ID No: 14)
14	g cga tat cta gaa tcc agc A ( SEQ ID No: 15)
15	gg gac aga gag acc agG ( SEQ ID No: 16)
16	cc caa aac ctg gag act G ( SEQ ID No: 17)
17	gtt tct gct gtt gct gct G ( SEQ ID No: 18)
18	ag acc tgg gtg gcc acT ( SEQ ID No: 19)
19	t gct gct g gct gct gcT ( SEQ ID No: 20)
20	c acc cgc agc gag gcA ( SEQ ID No: 21)
21	ctc ttc ctc tcc caa aac G ( SEQ ID No: 22)
22	gc tcc cag cat ttc tac taT ( SEQ ID No: 23)
23	cgg cga tat cta gaa tcc A ( SEQ ID No: 24)
24	g tca gct ctt ggg tcc G ( SEQ ID No: 25)
25	cc atg aag acc aag aca cT ( SEQ ID No: 26)
26	tgc caa gga gag gag caA ( SEQ ID No: 27)
27	gaa cta cgg cga tat cta G ( SEQ ID No: 28)
28	c cag cat ttc tac tac gat A ( SEQ ID No: 29)
29	gct gca gag ggt cca gG ( SEQ ID No: 30)
30	c tgg cgt cag gat ggg C ( SEQ ID No: 31)

Table 25-2

## Probe No.

## Base Sequence

31	ggc ttg cat tcc ctc cG (SEQ ID No: 32)
32	c cca gtt ggg acg agt gT (SEQ ID No: 33)
33	ct gct gct gct gct gcT (SEQ ID No: 34)
34	a gaa gat gtc ctg gga aaC (SEQ ID No: 35)
35	t gtg cag tca ggg ttt ctT (SEQ ID No: 36)
36	gcc tca gag ggc aac atC (SEQ ID No: 37)
37	ct gct gct gct gct gcT (SEQ ID No: 38)
38	ttc tat ccc cgg aat atc aT (SEQ ID No: 39)
39	gtt gct gct gct gct gcT (SEQ ID No: 40)
40	cag acc ttg gcc atg aac A (SEQ ID No: 41)
41	gg aat cac agc act cac G (SEQ ID No: 42)
42	a cgg cga tat cta aaa tcc A (SEQ ID No: 43)
43	ctc tcc caa aac ctg gag T (SEQ ID No: 44)
44	ttc ttg aag gaa gat gcc G (SEQ ID No: 45)
45	cat gaa gac aac agc acc aA (SEQ ID No: 46)
46	ggg ttt ctc gct gag gG (SEQ ID No: 47)
47	caa gga gag gag cag agT (SEQ ID No: 48)
48	g gcc acc agg att tgc G (SEQ ID No: 49)
49	c agg gct tct ggc ttc tG (SEQ ID No: 50)
50	ag aaa aca tca gct gca gaT (SEQ ID No: 51)
51	at caa cac cca gtt ggg aT (SEQ ID No: 52)

Table 26-1

Probe No.	Base Sequence
0	a gag acc agA gac ttg aca (SEQ ID No: 53)
1	ctg gag act Aag gaa tgg a (SEQ ID No: 54)
2	cga tat cta Aaa tcc ggc g (SEQ ID No: 55)
3	cta aaa tcc Ggc gta gtc c (SEQ ID No: 56)
4	c aca ctg aGc tgg cgt c (SEQ ID No: 57)
5	att att ttc taC gtc tgt tgt t (SEQ ID No: 58)
6	tg ctg tcc Ggg gat gga (SEQ ID No: 59)
7	acc cgc agT gag gcc tc (SEQ ID No: 60)
8	g agg aga aGa gtg ccc c (SEQ ID No: 61)
9	tg atg tca gCt ctt ggg tc (SEQ ID No: 62)
10	c ctg cgc tAt gac agg c (SEQ ID No: 63)
11	gaa tgg aca Gtg ccc cag (SEQ ID No: 64)
12	c aca ctg aCc tgg cgt c (SEQ ID No: 65)
13	gg att tgc cGa gga gag g (SEQ ID No: 66)
14	gaa tcc agc Ata gtc ctg a (SEQ ID No: 67)
15	a gag acc agG gac ttg ac (SEQ ID No: 68)
16	ctg gag act Gag gaa tgg (SEQ ID No: 69)
17	gtt gct gct G gct gct g (SEQ ID No: 70)
18	g gtg gcc acT agg att tg (SEQ ID No: 71)
19	gct gct g gct gct gcT a (SEQ ID No: 72)
20	agc gag gcA tca gag gg (SEQ ID No: 73)
21	tcc caa aac Gtg gag act g (SEQ ID No: 74)
22	at ttc tac taT gat ggg gag (SEQ ID No: 75)
23	cta gaa tcc Agc gta gtc c (SEQ ID No: 76)
24	t ggg tcc Gct ggc tcc (SEQ ID No: 77)
25	cc aag aca cTc tat cac gc (SEQ ID No: 78)
26	a gag gag caA agg ttc acc (SEQ ID No: 79)
27	cga tat cta Gaa tcc ggc g (SEQ ID No: 80)
28	tac tac gat Agg gag ctc t (SEQ ID No: 81)
29	g ggt cca gGg ctc gtg (SEQ ID No: 82)
30	cag gat ggg Cta tct ttg a (SEQ ID No: 83)

Table 26-2

Probe No.	Base Sequence
31	at tcc ctc cGg gag att ag (SEQ ID No: 84)
32	t gct gct gct gct gcT at (SEQ ID No: 85)
33	ct gct gct gcT att ttt gtt (SEQ ID No: 86)
34	c ctg gga aaC aag aca tgg (SEQ ID No: 87)
35	a ggg ttt ctT gct gag gta (SEQ ID No: 88)
36	g ggc aac atC acc gtg ac (SEQ ID No: 89)
37	gct gct gct gct gcT att (SEQ ID No: 90)
38	cgg aat atc aT a ctg acc tg (SEQ ID No: 91)
39	gcc atg aac Atc agg aat tt (SEQ ID No: 92)
40	gc act cac Gct gtg ccc (SEQ ID No: 93)
41	cta aaa tcc Agc gta gtc c (SEQ ID No: 94)
42	aac ctg gag Tct gag gaa t (SEQ ID No: 95)
43	gaa gat gcc Gtg aag acc (SEQ ID No: 96)
44	c agc acc aAg agc tcc c (SEQ ID No: 97)
45	c gct gag gGa cat ctg g (SEQ ID No: 98)
46	g gag cag agT ttc acc tg (SEQ ID No: 99)
47	agg att tgc Gaa gga gag g (SEQ ID No: 100)
48	ct ggc ttc tGt ccc tgg a (SEQ ID No: 101)
49	a gct gca gaT ggt cca ga (SEQ ID No: 102)
50	ca gtt ggg aTg agt gac c (SEQ ID No: 103)



Table 27-1

Allele Number	Probe Number for Detection				
	0	1	2	3	4
MICA*001	0	1	2	3	4
MICA*00201	5				
MICA*00202	6	7			
MICA*004	8	9			
MICA*005	10	11	12	13	
MICA*006	14				
MICA*00701	7				
MICA*00702	15	16			
MICA*00801	17	9			
MICA*00802	18	19			
MICA*00803	20				
MICA*00901	21	9			
MICA*00902	22				
MICA*010	23	13	9		
MICA*011	24				
MICA*01201	25				
MICA*01202	26				
MICA*013	6	27	13		
MICA*014	28	8			
MICA*015	28	29			
MICA*016	30	9			
MICA*017	31				
MICA*018	16				
MICA*019	32				
MICA*020	33				
MICA*021	34				
MICA*022	6	23	13		
MICA*023	6	17			
MICA*024	35	10	11	36	12
MICA*025	35	16			

Table 27-2

Allele Number	Probe Number for Detection		
MICA*026	7	37	
MICA*027	38	39	
MICA*028	27	17	
MICA*029	40		
MICA*030	41		
MICA*031	35		
MICA*032	25	42	8
MICA*033	43		
MICA*034	44	12	
MICA*035	6	38	
MICA*036	45		
MICA*037	38		
MICA*038	36		
MICA*039	30		
MICA*040	15		
MICA*041	46	5	
MICA*042	18		
MICA*043	47		
MICA*044	6	8	12
MICA*045	48		
MICA*046	49		
MICA*047	46	41	
MICA*048	50		
MICA*049	51		

Table 28-1

Allele Number		Probe Number for Detection				
		0	1	2	3	4
MICA*001	0					
MICA*00201	5					
MICA*00202	6		7			
MICA*004	8		9			
MICA*005	10		11	12	13	
MICA*006	14					
MICA*00701	7					
MICA*00702	15		16			
MICA*00801	17		9			
MICA*00802	18		19			
MICA*00803	20					
MICA*00901	21		9			
MICA*00902	22					
MICA*010	23		13	9		
MICA*011	24					
MICA*01201	25					
MICA*01202	26					
MICA*013	6		27	13		
MICA*014	28		8			
MICA*015	28		29			
MICA*016	30		9			
MICA*017	31					
MICA*018	16					
MICA*019	23		13	32		
MICA*020	33					

Table 28-2

Allele Number	Probe Number for Detection				
MICA*021	34				
MICA*022	6	23	13		
MICA*023	6	17			
MICA*024	35	10	11	36	12
MICA*025	35	16			
MICA*026	7	37			
MICA*027	38	32			
MICA*028	27	17			
MICA*029	39				
MICA*030	40				
MICA*031	35				
MICA*032	25	41	8		
MICA*033	42				
MICA*034	43	12			
MICA*035	6	38			
MICA*036	44				
MICA*037	38				
MICA*038	36				
MICA*039	30				
MICA*040	15				
MICA*041	45	5			
MICA*042	18				
MICA*043	46				
MICA*044	6	8	12		
MICA*045	47				
MICA*046	48				
MICA*047	45	40			
MICA*048	49				
MICA*049	50				

The present invention is not limited to the above embodiments and various changes and modifications can be made within the spirit and scope of the present invention. Therefore, to apprise the public of the scope of the present invention, the following claims are made.

This application claims priority from Japanese Patent Application Nos. 2003-430553 filed on December 25, 2003, 2003-430554 filed on December 25, 2003, 2003-430555 filed on December 25, 2003, 2003-430556 filed on December 25, 2003, 2003-430557 filed on December 25, 2003, 2003-430558 filed on December 25, 2003 and 2003-430559 filed on December 25, 2003, which are hereby incorporated by reference herein.